>SGPR397 SEQID 1

AGAGGAAGCATATGCACTGAAGAAAATATCCTATCAACTTAAGGTGGACCTGTGGCAGCCCAGCAGTATCTCCTA ATTGGTCCTGCCTTTTGTCAGTGGTTTGTAAAAGAAGCTCTTCTAACATATAAGAGTGACCCAGCCATGAGAAAA GCTGAATCATCTATATTCTATATCATGCCTGTGTTTAACGTCGATGGATACCATTTTAGTTGGACCAATGATCGA GCAACCGGGGCACACCTTTATAACAACCGCTATGCTGGTGATAAAGTGATAAGATTTATTCCCAAAACAGA ATTTTAAAGCTGGGCAGACGATCACGACTCAAAAGAGCTGTTTGGATAGACTGTGGTATTCATGCAAGAGAATGG IGTATCAGAGGGAACAGTTACTGATGTCCATATCCCCCAAAATGGTTCCCGAGCCCTGTTAGCCTTCTTACAGGA AGCCAACATCCAGTACAAGGTCCTCATAGAAGATCTTCAGAAACACTGGAGAAGGGAAGCAGCTTGCACACCC AGAGAAACCGAAGATCCCTCTCTGGATATAATTATGAAGTTTATCACTCCTTAGAAGAAATTCAAAATTGGATGCA GAAGTGGTGTGGTAAGTTTGGGACCAACTGGGATCCAGATCCAAAGGTTTCTGCAGGTTTTACTCTGCAAAATAT TTTGGAGAAAAACAAGGTCAAGGAACTCAAGGTTTCGCTGCCGTGGAGTGCCAATAGAAACTGGAAAGT ATGAAGTGTCTCGGGAAGCGCAGGGGCCAGGCAGCTGCTTTCCTGCCTCTTTGCTGGCTCTTTTGAAGATTC1 | CATCTGAATAAAACTCACTCAGGCCTCATTCACATGTTCTCTATTGGAAGATCATATGAGGGAAGATGTCTTTT GAGTCCAGAGGACTCTCATGGGAGACTCATGTTTTTCTGTATGTGA

>SGPR413 SEQID 2

FIG. 1A

GCCCAACACAGAGAGATTGTGGACAAGTCAGTGAGTCCATGGAGCCTGGAGACGTATTCCTATAACATATAC CATGTTTTGGGACGGATCTCAATCGAAATTTCAATGCATCTTGGTGTAGTATTGGTGCCTCTAGAAACTGCCAAGA AACATAGATGGTTATATCTACACTTGGACAACTGATCGTCTTTGGAGGAAATCCCGTTCACCCCATAATAATGGCA ATGAAGCČTCTGCTTGAAACCCTTTATCTTTTGGGGATGCTGGTTCCTGGAGGGCTGGGATATGATAGATCCTTA AGGATGATATTTTGTGCTTCCTGACCATGCACTCTTATGGGCAGTTAATTCTCACACCTTACGGCTACACAAAAA TAAATCAAGTAACCACCCAGAAATGATTCAAGTTGGACAGAAGGCAGCAAATGCATTGAAAGCAAAGTATGGAAC CCAGCCCACCTGTGAGGAGCCATGGAGGCTGTGTGTCAGTCCTGGATGATGTGTATGCGAAACACTGGCACT CACCCCATGGGAGAGATCTATGAGTGGATGAGAGAGATCAGTGAGAAGTACAAGGAAGTGGTGGTGACACAGCATTT CCTAGGAGTGACCTATGAGACCCACCCCATGTATTATCTGAAGATCAGCCAACCATCTGGTAATCCCAAGAAAAT CATTTGGATGGACTGTGGAATTCACGCCAGAGAATGGATTGCTCCTGCTTTTTGCCAATGGTTCGTCAAAGAAAT TCAAACATTCTGTGGGACAGGGCCAGTGTCTGAACCAGAGATAAAGCTGTTGCCAGCTTCATAGAGAGCAAGA
 IGGGATTCCCTTCTCATATACGTTTGAGCTGAGGGACAGTGGAACATATGGGTTTGTTCTGCCAGAAGCTCAGAT
TCTACAAAACCATAAAGACAACTCAAGTATACGCAAGCTCCTTAGGAACCTGGACTTCTATGTCCTTCCAGTTCTT CAATTATAGAGTTGGATCGAGTGCAGATATTTTATATGCCTCATCAGGGTCTTCAAGAGATTGGGCCCGAGACAT CGGACAGTGCTGGAAGGGTGACATCTGCCACTATGCTGCTGGGCCTGCTGGTGTCCTGCATGTCTCTTCTCTAA

>SGPR404 SEQID 3

AGCTGTATGCTGTGGAGATCTCAGATCACCCTGGGGAGCATGAAGTCGGTGAGCCCGAGTTCCACTACATCGCG GTTTCACTGCATCCACCAAGAACTGTATGGTTGGCTATGACATGGGGGCCACAAGGTGTGACTTCACACTTAGCA ACGGAGGACTTCCAGAAGGAGGAGGGCACTGTCAATGGGGCCTCCTGGCACACCGTCGCTGGAAGTCTGAACG ATCACCGCCGGAACGAGATGACCACCACTGATGACCTGGATTTTAAGCACCACAATTATAAGGAAATGCGCCAGI GAATTGACATCAACAACAACTTTCCTGATTTAAACACGCTGCTCTGGGAGGCAGAGGATCGACAGAATGTCCCCA AGAGCAGTCATAGCCTGGATGGAAAAAATCCCTTTTGTGCTGGGCGGCGACCTGCAGGGCGGCGGCGAGCTGGTGG TGGCGTACCCCTACGACCTGGTGCGGTCCCCCTGGAAGACGCAGGAACACACCCCCCACCCCGGACGACACAC GGCTTGGTGAGAGATTCACATGGAAAAGGAATCCCAAACGCCATTATCTCCGTAGAAGGCATTAACCATGACATC AAACCAACATGGCCAGGATCCGAGAGATCATGGAGAGTTTGGGAAGCAGCCCGTCAGCCTGCCAGGCGG ATGCCTACGAGAGGCCTACGAAGGGGGCTCGGAGCTGGGAGCTGGTCCCTGGGACGCTGGACCCACGATG AGCTGCCCGAGGAGTGGGAGAATAACCGGGAATCTCTGATCGTGTTCATGGAGCAGGTTCATCGTGGCATTAAA GTTCCGCTGGCTGGCCTACTCCTATGCCTCCACACACCGCCTCATGACAGACGCCCGGAGGAGGGGTGTGCCAC CGAACAGCCAACGATGGGGGATTACTGGCGCCTCCTGAACCCTGGAGGTATGTGGTCACAGCAAAGGCCGAAG ATGGTGAGCAATGACAGCCACACGTGGGTCACTGTTAAGAATGGATCTGGAGACATGATATTTGAGGGAAACAG 1 GAGAAGGAGATCCCTGTTCTCAATGAGCTACCCGTCCCCATGGTGGCCCGCTACATCCGCATAAACCCTCAGT SCTGGTTTGATAATGGGAGCATCTGCATGAGAATGGAGATCCTGGGCTGCCCACTGCCAGATCCTAATAATTATT IGATGAAAGTTGTGAATGAAATGTCCCCAATATCACCAGAATTTACAACATTGGAAAAAGCCACCAGGGCCTGA GGAAAGTTCCCAATCACTATATTGCAATCCCTGAGTGGTTTCTGTCGGAAAATGCCACGGTGGCTGCCGAGACC ATTTCAGCTACCTTCATACAAACTGCTTCGAACTGTCCATCTACGTGGGCTGTGATAAATACCCACATGAGAGCC GCTGAAGCTGCGGGGGGGGGAAGAGACGACAGCGTGGGTGA

>SGPR536_1_SEQID_4

CAGGAGCAGGACCTGTGCTGCCGCGGCCGTGCCGACGACTGTGCCCTGCCCTACCTGGGCGCCATCTGTTACT GTGACCTCTTCTGCAACCGCACGGTCTCCGACTGCTGCCCTGACTTCTGGGACTTCTGCCTCGGCGTGCCACCC GTCGTGGGCGCCGGGAGCTAGCACCGGGTCTGCACCTGCGGGGCATCCGGGACGCGGGAGGCCGGTACTGC AACCGTTGCACCTGCCAGGAGAACAGGCAGTGGCAGTGTGACCAAGAACCATGCCTGGTGGATCCAGACATGA TCAAAGCCATCAACCAGGGCAACTATGGCTGGCAGGCTGGGAACCACAGCGCCTTCTGGGGCATGACCCTGGA

CCTTCTCGGGCCGTGAACGAGACGAGGCTGGCCCTGCGCCCCCTGTATGATGCACAGCCGAGCCATGGGTCG TCTACCGCCTCGGCTCCAACGACAAGGAGATCATGAAGGAGCTGATGGAGAATGGCCCTGTCCAAGCCCTCATG AGAGATACCGCCGGCATGGGACCCACTCAGTCAAGATCACAGGATGGGGGAGAGGAGACGCTGCCAGATGGAAG GGGCAAGCGCCAGGGCCACTGCCCCAACAGCTATGTTAATAACAATGACATCTACCAGGTCACTCTG CTTGACCAAGGCAACTGTGCAGGCTCCTGGGCCTTCTCCACAGCAGCTGTGGCATCCGATCGTGTCTCAATCCA GCGCGGCGTCAATGAGTGCGACATCGAGAGCTTCGTGCTGGGCGTCTGGGGCCGCGTGGGGCATGGAGGACAT GAGGTGCATGAGGACTTCTTCCTATACAAGGGAGGCATCTACAGCCACACGCCAGTGAGCCTTGGGAGGCCAG GCCGCGGTGGGCGTCTCGATGGTGCCTGGTGGTTCCTGCGTCGCCGAGGGGGTGGTGTCTGACCACTGCTACC GCTGAACCCAGGGGAGGTGCTTCCCACAGCCTTCGAGGCCTCTGAGAAGTGGCCCAACCTGATTCATGAGCCT **GGGTCATCACTGA**

>SGPR414_SEQID_5

GACCAAATTTACATATTGAGATTATCAAACAGTGCCAAGTGATTTTTGAATTTTTGGCAGCAGAAGGGGCGACTGAG IACTCAACATATTGACTGTATTTGGGCTGCAGCACAGTTGAAACATTGTAGTCGGTATATACATGACTTATTTCCTT AAGGAATATAAGCATTTGGAGATTTTTAATCAAGTAGTGTGTGCACTTATTAACTTAGTGATTGCCCAAGTTCAAGT <u> ATGTGCGĀGAACTĒCGCAGACCTGGTGGAGGTGTTAAATGAAATATCAGATGTAGAAGGTGGTGATGGACTGCA</u> GCTCAGAAAGGAACATACTCCAAAATATTTACTTACATCAATTCCTGGACACAGAGGCAATGTCTATGCTGCTTC GCTCCGGGGACCAGCTTTGTAAACATTGTACTACCATTAACATAGATTCCACGTGGCAAGATGAGAGTAATCAAGC AACTCTACAAGAATTTGTAATCTGACTGAGGAGTTCTTCAAAGAGTTCTGATCTTTAGTTTATGGAGTACAG ATGAGAAGGAAAAACTCTTACTATGTGGCAAAAATTTTTCAAATTCAGTTTCCCTTATATACTGCTTACAAGCAT CTTTGAATATGGAACTCCTGAAACTTTGCCATTTCTTATAGCACATGCGTTTATTACAGTTGTGTCTAATATTAGAA TGGATACAACATTATGCTTTGATAAAGAAAGCCTAGATCTTGCATTTAAGTACTTTATGTCACCTACTTTGACTATG CGGACACAGAAACGTCCATTGCAAAAGAACTTGCAGACTGGCTTATTAGCAACAATGTGGTGGAGCATATATTTG AGGTTGGCTGGATTGAGTCAGATAACAAATCAACTCCATACCTTCAATGATGTGTGCAATAATGAATCATTAGTAT TAGAAGTACCATTGCATTTGCTTCGTTATGTATGTTTTGTGGGAAAAATGGCCTTTCTCTCATGAAGGATTG CTCGGATCAGGAGTTACGACAGAGTGCAGCTCGTAACATGGCTGACTTAATGTGGAGCACAGTCAAAGAACCAT TATGGCTACATATTCCCGCTGTCATGCAGCACATTATACCTTTTAGGACCTATGTTATTAGGTATTTATGCAAGCT

GAGCAAGTTGACATCTTATGGCATTGTTTAGTAGAAGATTCTGAATGTTATGATGATGCACTCCATTGGTTTTTAAA AGCAAGAATTTATTAGTAAGTGCATGGAGAGTCTTATGATAGCTTCTAGCAGTCTTGAACAGGAATCACACTCAAG GCCTTGAAAACTTGGGAAACAACAGATCAGTAGTAATTTCACTTCGTCTTCCAAAAACTATTTGGTACTTTTCAG GCTAAAACCTGAAACAATTAGCATGACTGGCTTAAACCTGTTTCAGCATCTCTGTAACTTGGCTCGATTGGCTACC GCTAAGGGTTGTATGCCAGCCAGCTGGACTTCCTGACAAGATGACTATTGAAATGTATCCTAGTGACCAGGTAGC ICTGGTGATGTCAGTCGAGCAGCTATCCAGTATATTAACTCCTATTATATTAATGGTAAAAACAGGTTTGGAGAAGG AATCATTGCCATCAGTAGATAATCGAATGCGAATGCTGGATGCTTGTTCACACTCTGAAGACCCAGAACATGATAT GATTAATTCACATGCGGAACTGACATCTCACCTCCAACAACATCTTCCCAATTTAGCTTCCATTTACCATGAACAT CTTAGTCAAGGACCTGTAGATTTGGATAAACATCAATTCAACAGACATGATTACAGACATTACAGACATTAGATTTGGATAATGTTTG ICTGAGACAGTGGCAAATTGAAGGCACTGGTATTAGTAGTCATTTGAAAGCACTGAGTGACAAACAGTCTCTGCC CCATCACCACCACCACGATGGGCATATGGTTGATGTTGCTAAGTGCAGATGATGTCAGTTGTAGTAGCTCCCA CAGTAGTGGGCATAGTGATGGATCTAGCAATGAGGTTAATTCTAGCCACGCAAGCCAGTCAGCTGGGAGCCTG CATAATCCTCCCAAAAGCAGTTGTGGTACAGATCTTCGGAATAGAAAGTTAGAGAGTCAAGCAGGCATTTGCCTG CAGTTTGGGAGCAGTTACGATACACACTGGATAACAATGTGGGCAGAAAAAGAACTGAACATGATGAAGAAGTTTTC AAGTTCAAGTTCGTCTTCAATTCTTGACTTGTGTATTTTCAACTCTGGGATCACCTGATCATTTCAGGTTAAGTTTA TCTCATGGTTATAGAAAGAGGACTCCTTATGCTGAAGACACATCTGGAAGCGTTTAGGAGAAGGTTTGCATATCA CACTCATCAAGAATTTGGATCCCGTACCACTTAGACATCTACTTAATCTGGTCTCAGCTCTTGAGCCAAGTGTTCA TTCAGGGGAAATGAATGCTACTCATATAGCACAAGGGTCTCAGGAGTCTTGTATCACACGAACTGGGGACTTCCT **GGTTAGTGCAAAATCAGAAAAAAAAAAATATGGCTGATTTTGATGGTGAAGAAATCTGGATGTGAAGAGAGGAGCTAGTTCA** TCAAGTTCGAAGTAAAGATCAACATGCTATGGGTATGGAAACCTACAAACATCTTTTCCTGGAGAAGATGCCCCA AGTGCCTATGATGGTTGTTCAAATTCTGAGCTGTGTGGTATGGACCAATTTTGGGGGCATTGCTTTAAGAGCACAA TAGAAGAACAGCTCCATCACCTTGGTCACCTGCAGCTAGTCCTCAAAGCAGTGATAATAGCGATACACATCAAA FACT GAACAGACACTGTACTTGGCATCCATGTTAATTAAAGCACTGTGGAATAACGCACTAGCAGCTAAGGCTCA GTGGAGGTAGTGACATTGAAATGGATGAGCAACTTATTAATAGAACCAAACATGTGCAACAACGACTTTCAGACA GGGGACTCCCAAGGCACGTCAGAAAGAAATGGGACAAGCGGGAACAGGAAAGGAACTGGTTTTAACACTG CAAGAAAGGAAATACTTTGTTGTGGGATATAGTCCAAGATGAAGATGCAGTTAATCTTTCTGAAGGATTAATAAAT GCAGTGAGGTACAGTCAGAAGACATTGCAGATATTGAAGCCCTCAAAGAGGAAGATGAAGACGATGATGATGATGGT TGGGGAGACTATTGGGAATGAATTATTTAATTGTCGACAATTTATTGGTCCACAGCATCACCACCACCACCACCACCA GAAGCAGAGAAACTTCTTTGTTCGTTAGTATGTTGGTTTACAGATAGACAAATTCGAATGAGATTCATTGAAGGTT CAGAGGAATCCATGCAGGGAAGTTCTGACGAAACTGCCAACAGTGGTGAAGATGGAAGCAGTGGTCCTGGTAG

AATGTAGAAGATGATGGGCTTACAGGACTCCTAAGGCTTGCAACAAGTGTTGTTAAACACAAACCACCCTTTAAAT ICTACTATTCAGCAACTTTATATGATACCTGAGGCAAGACAGGCTGTCTTCACTGCCAAGTATTCAGAGGATATGA AGATTGGCTAAAGAAATTGACTCTCCTTATTCCTGAGACTGCAGTTCGTCATGAATCATGCAGTGGTCTCTATAAG AATTTCTTCCTGATGCTCAAGCACTCAAACCTATTAGGATAGATGATTATGAGGAAGAACCAATATTAAAAACCAGG CTGGCCTCATGAAGATGTCCGTGCTGAATGTAGATTTGTTGGCCTTACTAACCTTGGAGCTACTTGTTACTTAGCT AGCACAAGACCACTCTTCTGGAGCTTCAGAAAATGTTTACATATTTAATGGAGAGTGAATGCAAAGCATATAATCC CAATGTGGCTCTTGGTGAGTTGGGCTCATTGCTGTTCTTTAGTGAAATCTAGCCTTGCTGATAGCGATCATTTACA CTCGACTTAGATGCCTTGGCAAGACATTTGGCTGACTGTATTCGAAGTAGGGAGATCCTTGATCATCAGGATGGT TTTCAAGGGAAGGACAGGAATTTTTGAGAGATATCTTCAATCTCCTGTTTTTGTTGCCAAGTCTAAAGGACCGACA GCCAAGTGGCTGATATGAAGAACATTTATGAATCTCTTGATGAAGTTACTATAAAAGACACTTTGGAAGGTGATAA CATGTATACTTGTTCTCAATGTGGGAAGAAGTACGAGCTGAAAAAAGGGGCATGTTTAAGAAATTGCCTCGCATI AGAAGTCAGTGATCATTCAAAAGACTCAGAGGCTATGAATATGACTTGATAGGAGTGACTGTTCACACAGGAAC CTTTTAATGATGCTGAGGTAAAACCTTTTGATTCTGCTCAACTTGCATCTGAATGTTTTGGTGGAGAGATGATGACGA CCAAGACCTATGATTCTGTTACAGATAAATTTATGGACTTCTCTTTTGAAAAGACACACAGTGCATATATGCTGTTT TGAGAACTACAGGCTAATACACAACTGGGTTATGGCACAACACATGCAGTCCCATGCACCTTATAAATGGGATTA TTNAGTTTCAATACTATGAGATACACATTTAATATGGTCACGATGATGAAGAGAAAGAGAAAGTGAATACACACTTTTCCTT CCCATTACGTTTGGACATGACGCCCTATACAGAGATTTTCTTATGGGAAAGAGGGGAAAGAAGAAGAAGGATTTAA **GGCAGATGGTGGACACTATTATAGCTTTATCAGAGATATAGTAAATCCCCATGCTTATAAAAACAATAAATGGTAT** ACACGAGTTAACAACAGATTATGATGAAAAAGCACTTCATGAGCTTGGTTTTAAGGATATGCAGATGGTATTTGTA CATGCAGAAAATCTGTCTAGGCGGGTCTGGGAGCTACTGATGCTTCTTCCTACATGTCCTAATATGTTGATGGCA GAGTTTGGTCAAAGCAACCGAAAAGGAGAGTTTCCTGGAGGCCTCATGGGACCTGTCAGGATGATTTCATCTGG ACAGCCAAAGTGCAAATCACATTCTTCAAGAGCTGCCGCTTACGATTTGTTAGTAGAGATGGTAAAAGGGGTCTGT TCTTTGGGTGCACCAAGGAGAGAGGGGAAAGGGGGAAGGTGTTCAGCTGCCAGCATCTTGCCTCCCACCCCTC TTCCAGAATATCTCAGATGAGCAGAGTTTTAAAGCTCAGTCTGATCACAGGTCTAGACATGAAGTTTCACATTATT TAGACCTTTCTGTAAAACATACACCATGGATAAGCAGCCTCTGAATACTGGGGAACAGAAAGATATGACAGAGATT GTAATTACAAACAATGTTGTATCCTTGGATTGTGAACATGTTAGTCAAACTGCTGAAGAGTTTTATACTGTGAGGT GCATCATTTAAACCACCCTCAGGAAAAGTGGCAGTGGATGATAGTGAGAGCTTACGATGTGAAGAACTTCATCT AGAAGGACAACATTCCAATGCTTTTGCTTTTACAAGAGCCTCATTTAACTACTCTTTTTGATTTATTAGAGATGCTT ITTTACTGATCTAATTACCAAAATCGAAGAAATGTCTCCCGAACTGAAAAATACCGTCAAAAGTTTATTTGGAGGT

GATTGATTCTAATGACAGAGTCTTTCAACACTTTGCACATGATGTATCACGAAGCTACAGCTTGCCATGTGACTGG GACTGACATGGCAGCATTAACAGGAGGAAAGGGATTTCCCTTCTTGTTTCAACATATTCGTGATGGCATCAATATA TCTCAGTGTCTAGATTGGTTGGCAGTGCAGACACCCCGAAATAAACTGGCACACAGGCTGGGTCTTACAGAATATG TCCGCCTTATTGTTCAGAACCCAGTGGTAACCAAGAACATTGCCTTCAATTACATCCTTGCTGACCATGATGATCA AATACCCTGGAGCAGTAGAAGAACTGTTTAACCTGATGCAGCTGTTTATAGCTCAGAGGCCAGATATGAGAAAG AGATITAGTAGAACTICTGTCAATATITCTTTCGGTTTTGAAGTCTACACGCCCTTATCTTCAGAGAAAAGATGTGA CTGCATTCACACGACAACTGGCTTCTCACCAGAACATCCAGTGGGCCTTTAAGAATCTTACACCACATGCCAGGC ITAAGTGCCCTAATCAAATTGTGAGACAGATGTTTCAGCGTTTGTGTATCCATGTGATTCAGAGGCTGAGACCTGT GCATGCTCATCTCTATTTGCAGCCAGGAATGGAAGATGGGTCAGATGATATGGATACCTCAGTAGAAGATATTGG CCACTCAGTCCAGACACAACAGTAGTCCTACATCAGGTCTACAACGTGCTCCTTGGTTTGCTCTCAAGAGCCAAA GCAATAGCTACAAATCACAATAAACAGGCTTTGCTTTTGGTACAATGTCTGTGCTGACTGTCCAGAGAATA TACAAACGCATGGAACCAGAGGAAGAAATGGCAGAGAATACAAATTTGATGTTTCGTCAGAGTTACTAGAGTGG GTCCTAGAGACATTTATTCATTCTAAAGAAAAGCCCACGATGCTTCAGTGGATTGAACTGTTGACGAAACAGTTTA ATAATAGTCAGGCAGCTTGTGAGTGGTTTTTAGATCGTATGGCTGATGACGACTGGTGGCCAATGCAGATACTAA GCAAGCTATATCTACAATGGTACATTTTTACATGGGAACAAAAGGACCTGAAAATCCTCAAGTTGAAGTGTTATCA CCCTTGAAAAGATGATAGCTTTTAGTTGCTCTTTTGGTTGAACAGTCTCGATCAGAAAGGCATTTGACATTATCACA GGTGGACCTCCAGGAATGCCTCCCTTTGCATCTTATATTCTGCAGAGGATATGGGAGGTGATTGAATACAATCCT GAAAACTGGGTCGAGCGGTTTCTTTTGGCTCACAATTATCCTAGAGTGAGGACTTCTGCAGCTTATCTTCTGGTG CTTTATGTTGATGCTGCTGTTCATGGCACTACAAAGCTAGTGCCCTATTTTAGCTTTATGACTTACTGTTTAATTTC CAAAACTGAGAAGCTGATGTTTTCCACATATTTCATGGATTTGTGGAACCTTTTCCAGCCTAAACTTTCTGAGCCA GGATGTGGTGCTTTTTAACCGTGGGATGCTGCCAGCGTACTATGGCATTCTGAGGCTCTGCTGTGAGCAGTCTC AACAAGCATTAATCCAGTGGCAGGGAGTTGAATTTGCCCATAAACTGTTAACTCTTCTTAATTCCTATAGTCC AGACAAACTTGTAATCTGATTTTCAGCCTGTGTCGATACAATAATCGACTTGCAGAACATATTGTATCTATGCTTTT TCCCTTATACCAAGCAATTCATTCCGTCAGATGTTCCGGTCAACAAGGTCTTTGCACATCCCAACCCGTGACCTT GTGTAGTTGTATTCCCAGTACATTACCAGATCCTAAAGCTGTGTCCTTAATGACAGCAAAGTTAAGCACTTCCTTT

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>SGPR430 SEQID 6

CTCAGTCTGCTATGTCAAAAAACTGCATCAAGCTTTTGTGTGAAGATCCTGTTTTCGCAGAATATATTAAATGTATC GCTTTGCTCCTGTCAGTACACACTCCCAAACAGTTAAACCCAGCTCTAATTCCAACTCTGCAAGAGCTTTTAAGCA CTAATGGATGAAAGAACTTTTTTAAACAACAATTGTCTACACGTTCATGACACATTTCCTTCTAAAGGTTCAAAG AAGGAGCAACTCCCATTAAAAGGCGGCGTGTTAGCAGTGATGAGGAGCACACTGTAGACAGCTGCATCAGTGAC TATCTCTGCAGTTCTGTCTGACTTAGCTGACTTGAGAAGCTGTGATGGCCAAGCTTTGCCCTCCCAGGACCTTGA GGTTGCTTTATCTCTCAGTTGTGGCCATTCCAGAGGACTCTTTAGTCATATGCAGCAACATGACATTTTAGATACC CTGGTTCCCTTTCTACAACACAACCATTGTACTTACCATCACAGTAATATACCAATGTCTCTTGGACCTTATTTCCC TTATCATCAGTTCATCCATCTATTATGCCGAGTTGCAATCAACTGTGAAAAATTTACTGAAACATTAGTTAAGCTGA TTGTCGAGAAAATATCAAGCTAATAGGAGGGAAAAGCAATATTCGGCCTCCGCGCCCTGAACTCAATATGTGCCT TCAAGTGTTTTCTGAAGCAAACTGTGCCAATTTGATCAGCACTCTTATTACAAACTTGATAAGCCAGTATCAGAAC CTACAGTCTGATTTCTCCAACCGAGTTGAAATTTCCAAAGCAAGTGCTTCTTTAAATGGGGACCTGAGGGCACTC ATGAAAACAGAAACCAGGGAGGTCCTGACCCCAACGAGCACTTCTGACAATGAGACCAGAGACTCCTCAATTATT GATCCAGGAACTGAGCAAGATCTTCCTTCCCTGAAAATAGTTCTGTTAAAGAATACCGAATGGAAGAAGAATTCCATCTT GTAAAGAATTTAAAGACCTCCACTGTTCCAAGGATTCTACCCTAGCTGAGGAAGAATCTGAGTTCCCTTCTACTTC CTGTGTAGGACCATTGAATCTACAATCCATGTCGTCACAAGGATATCTGGCAAAGGAAACCAAGCTGCTTGA GTGTCCTAGTTGCCTATGAAGGTTTGCCACTTCATCTTGCACTGTTCCCCAAACTTTGGACTGAGCTATGCCAGA CTTGCCCACAATGGTGGAAACCAGTAAGGGCAAAGATGACGTTTATGATCGTATGCTGCTAGACTACTTCTTTC CGTTTTCAGAAGACATGTCAAATATCAGGTCACAGCATGCAGAAGAACAGTCCAACAATGGTAGATATGACGATT TCCAGAACTTAGAAATGCCTGTATAGATGTCCTCAAGGAACTTGTACTTTTGAGTCCCCATGATTTTCTTCATACT

GACCAAAAAGGATTTACTCAAGAAGGTTAAAAATGCCATTTCAGCTACAGCAGAGAGATTCTCTGGTTATATGCAG **GGGCAGCAAGTCATTCCAAGATACCTGACCCTGTCATCTCATTGCACTGAAAATACAAAACCACCTTTTACCC** ATGAGAATAAAGAAAACAAAACTCCAGAAGGATCTCAGGGAGAGAGTTGATTGGCTCCAGCAGTATGATATGGAGC AAAATCCAAGAGGCTGCCGTGCAGAGTGATCGAGATCGGAGTGGCTACATCTTTTATATGCACAAGGAGATC CTTTAGGGCCGAAGAACTGGAGTATTCTTGTGAGAAGTGTGGTGGGAAGTGTGCTCTTGTCAGGCACAAATTTAA ITCTGATGAGGACTCTGGAAATGAGGATGTTTTTGATATGGAGTACACAGAAGCTGAAGCTGAGGAACTGAAAAG GAAGTTTGGGATTTCTTCCTCAGCCAGTTCCTCTTTCTGTTAAAAAACTGAGGTGTAACCAGGATTACACTGGCTG AGACTGAACCTGTTTCTGGAGAAAAAATTCACCAGATATTTCAGCTACCAGAGCATACACTTGCCCTGTTATTAC ATGATAAGCCAACTAGCAGCCCAGATACCGGATTTGCAGAAGATGATATTCAAGAAATGCCAGAAAATCCAGACA CTATGGAAACTGAGAAGCCCAAAACAATCACAGAGCTGGATCCTGCCAGTTTTACTGAGATAACTAAAGACTGTG GATCCAAGGAACACTCTTCTGGTGGCACTAACTTAGACAGGACTAATGTTTCAAGCCAGACTCCCTCTGCCAAAA GAATAAACCAAGAGTGCCCCTTTCCTCTCACCAACAGCAGCAACTGCAGGGCTTCTCCAATTTGGGAAATACCTG TTGGTTGGAGTGCACATATGGCAATGTCTAGACCATTGAAAGCCTCTCAAATGGTGAATTCCTGCATCACCAGCC CTTCTACACCTTCAAAGAAATTCACCTTCAAATCCAAGAGCTCCTTGGCTTTATGCCTTGATTCAGACAGTGAGGA TGAGCTAAAACGTTCTGTGGCCCTCAGCCAGAGACTTTGTGAAATGTTAGGCAACGAACAGCAGCAGGAAGACC GTGAAAGGGAAGAGCAAGAGCTTCAGCAGGCACTGGCTCAGAGCCTTCAAGAGCAAGAGGCTTGGGAACAGAA AGAAGATGATGACCTCAAAAGAGCTACCGAGTTAAGTCTTCAAGAGTTTAACAACTCCTTTGTGGATGCATTGGG TTGATGAGCTGCTGGAAACAGAAAGAACTCTCAGTCACTTAGCACGGAAGTGGGGAAGACTACCCGTCAGGC TAATTTGGAGTTTGAGGTTCAGCACTCCATCATTTGTAAAGCATGTGGAGAGATTATCCCCAAAAGAGAACAGTTT CAGGCTTCCTAGGGTCCTCATTCTCCATTTGAAACGATATAGCTTCAATGTGGCTCTCTCGCTTAACAATAAGATT AACAGTCAGAAGAGAATAGGACATCAGGTGGGCTTTTACCTTTACAGTCATCCTTTTATGGTAGCAGAGCTG TGGAAAAAGATTCAAAATTATGCCCAATAGAGCCTGACAAGTCTGAATTGGAAAACTCAGGATTTGACAGAATGA CTATATGAATGCTATTCTACAATCTCTATTTTCACTCCAGTCATTTGCAAATGACTTGCTTAAACAAGGTATCCCAT

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GGGAGGGAGAGCAGTGTGAACGGGGAAAGCCACATACCAGGAATCACCACCCTCACGGACTGCTTGCGGAGGT GGCGCAAGATCACTACATACATTTCCTTTCGTGGAGCTGGATATGACGCCGTTTATGGCCTCAAGTAAAGAGA GCAGAATGAATGGACAATTGCAGCTGCCAACCAATAGTGGAAACAACGAAAATAAGTATTCCTTGTTTGCTGTGG AAACAGCTCACAATGAATAAATTACCTGTCGTTGCCTGTTTTCATTTCAAACGGTTTGAACATTCAGCGAAACAGA TTAATCACCAAGGAACCTTGGAGAGTGGCCACTATACCAGCTTCATCCGGCACCACAAGGACCAGTGGTTCAAG TGTGATGATGCCGTCATCACTAAGGCCAGTATTAAGGACGTACTGGACAGTGAAGGGTATTTACTGTTCTATCAC CCACGATAGACCCATGCTGGGACATTAGTTTGGACTTGCCTGGCTCTTGCACCTCCTTCTGGCCCATGAGCCCA TTACGAGGCCAGAGCACTTAGGAAGCAGTGCCAAAATCAAATGTGGTAGTTGCCAAAGCTACCAGGAATCTACC

AAATACAGAAGGGAGGAAAATGCAGAAATTTATCTGTAAGAGGAATTACAAATTTAGGAAATACTTGCTTTTTAAT GCTTTGTCAGAAGGCACCTCGATTTAAAGATTTCCAGCAACAGGACAGTCAGGAGCTTCTTCATTATCTTCTGGAT AGAGAGCAATAGCTGAGAATCTGTGGTCAGTTTGCTCAGAATGTTTAGAAGAAGAAGAAGATTCTATGATGGGCAGC AAGAGTCAACTAATTCATGACCGAAAATGTATTAGAAAATTGTCATCTGGAGAAACTGTCACATACCAGAAAATG FACTGATGACAGTGAAAAAAGAAGCCAGCCATTCTGAAAGCAATGTTGATGCTGACAGTGAGAGCCTTCAGAATCTGA GAAGACTCTTCAGATGATATTGCTGTAGGTTTAACTTGCCAACATGTAAGTCATGCTATCAGCGTGAATCATGTAA AAGTECTTCAAAGCAGACTGGGCTGTTCAGATCCAGTAGTGGATCCGGTGTGCAGCCCAGATGGACCCCTTTACC TCAAGATTTTTCCTTCCTCAGACTCTCAGCTGGACCCATTGGTGGTGGAACTTTCAAGGCCTGGACCACTGACCT CAGCCTTGTTCCTGTTCACAGCATGAAGGAGTCTGAAAAAGGACCACTTTCTCCTAAAGGTTCTTTTAATCA -CGATACAGTGGCAATGTTACTATAGAAAATATTCATCAACCTAGAGCTGCCAAGAAGCATTCTTCATCTAAAGAT ATGCGGGTGAAAGATCCAACTAAAGCTTTACCTGAGAAAGCCAAAAGAAGTAAAAGGCCTACTGTACCTCATGAT TAGTACTTACTTCTGATATTTGGTTGTGCCTCAAGTGTGGCTTCCAGGGATGTGGTAAAAACTCAGAAAGCCAAC AAACATGCTTCTAAAACACAAACAAGTGCATTTTCTAGAATCATGAAACTTTGTGAAGAAAAATGTGAAACAGATG GTGTTATGAATGTGAAAAAATTATCAACGCATTGTAATAAGAAGGTTTTGGCTCAGATAGTTGATTTTCTCCAG GCAGTGAGGACAGAAGAAACAAAGCGAATACAAGCTAGCATTCTAAAAGCATTTAACAACCCAACTACTAAAACT CTCTGTCAGCAGGTAAACTGCTGTACACCAAGGAGACTGACAGTGGTGATAAGGAAATGGCAGAAGGTATTTCT >SGPR495 SEQID 8

GAACTTCGTTTGAGCAGCACTGTAACTGGGGATCAAGATTTTGACAGAGAAAATCAGCCACTAAATATTTCAAATA AGAAGGAGTTTATACTAATGCCAGGAAGCAATTGCTCATTTCTGCTGTTCCAGCTGTCCTAATTCTCCACCTGAAA CATAACACTAAAAAGAAAAATGTGCCTGGTTTGAAAGCGGCTGATAGTGAATCAGCAGGCCAGTGGGTCCATGTT -ATAACTACTTCTAAAGAATGTTCAATTCAGTCCTGTCTACCAGTTTACATCTATGGAATTACTAATGGGAATA AGATTTCATCAGGCTGGCTTGAGTCTTCGTAAAGTAAACAGACATGTAGATTTTCCACTTATGCTCGATTTAGCAC CATTCTGCTCTGCTACTTGTAAGAATGCAAGTGTGGGAGATAAAGTTCTCTACGGTCTCTATGGCATAGTGGAAC ATTTATGTTTTTAGAGGGAAGCATTTGAGGTCTTATAGTCCCCAAAATGCTTTTCAGACCCTTTCTCAGAGCTA ATAGTGGCTCGATGAGAGGCCACTACACTGCTTATGTGAAAGTGAGAACACCCTCCAGGAAATTATCGGAA

>SGPR407 SEQID 9

TGGCTGGTGTTGGTGCCTCTGAAACTGACAGGAAGTCCCCACAGATGGAGGGCCCAGGAAGAGGGGCCTGGCCA GAACCACTCAGGCAGCATCCACTGTGGCCACTACACACCCTGTGCCAGTGCCAGACTGGTTGGCACGTTTACA ATGGAGTATCCAGTCCCATACTTTAGATCCCCGAACAGGACTCTGATCCCAGAGAGAATTTGGTCAAACCCATTA ATGCCCTTTGTGATACCGAGCAAAACCTTGCCATGGGACCCACTGGAACTCAAGATTTGTTATCAGCAAAATCGC CCATATCCCTCCCCGGACCCATCAAACTTTCCTACCTTCTTACGCTGTCTGAATGCTTTCTCTGCAGCTGTCTTCT TATGGGAGCGGCTACACCTTGAAATCAACCACGGAGGCCGCGGGGCTCCACCAGTCCCTGCCCATGGTCCAGC FCCCTCTCCACCCACCAAGGGGGAGTGCTCTGCTAAAAGAGTCTGAGTTAAATGATGATGACTGGGCCAACCTA ATGTGGAAGCGTTATCTGGAAGAACAAGAGGACAGCAAGATGGTGGATCTGTTGTGGGCCAGATGAAAAGTTA AGAGTTAGAGAATGCCTCAGGGACTTTGCCAGTGACAAAGTCGGAAGTCCTGTCTACCAGCTGTGTGCCCTTTG CTAGACATTCTGTTCAAAGCTTCCACAGATATTATTTTAATCATGACTCCAGCTCTGGGAACAAATGGAGGA AGTTACCAGAACCTGGAGGTTTGGAAAAGAAACATGAAGAGCTGAGACTCAGACCTCTGAAGGAGGAGTACCAT GCTGCAGCTGGTGTCTCCAAAGGGTCACCATGAGGCGGGTTATGGGTGTGCAGGACAAAGCTGGAAACAGGAA GCTTGCAGACGGCCACCTTCTCACTCTTCACACAGTCCTTCTGGGCATGTCTGGATCCTGATCTTTC CCCAAGAAAGGATTTGCTGGGGGCAAGGTGTCTCTGCGGGATTGTTTAAGCCTTTTCACCAAGGAAGAAGAGGCT ATGACTCTTGTGTCTCCCCTAAACACGCTGCGGGACACAGAAGGAATAGAACTCACAGTTATGAAGGCTCTAGTI CCAGATGCTGCTGCTGGGGCAAAGACCTGTGATAGGTGATACAGTCAGCAACAGCCAGACAACTAGGGACAAG ATCTCCCACAGCCCTCATGGCATAAGCCCGAGGGCTTAAAGCCAGCAGGATACCCAAGAGTTCCTGACATTCCT

AGTGGGATGGAGCCACAGAAGTCTGCACCATTTGCAGCAGGGAAGGGTCTGGCCCCTCCTCTTCCTGTGTGCAA AACTCTTCCAGCCTGATGCTCATGGATTTCTAGTGAAGAAGTTCATGCTCCAACAAGGGGCATCGTGTTTATCAT ATGGCCCAGAGTTTTTACTGGGGTTTTCCTGCTGGAGGGAACTTGTCTGCTTTAGAAATGCTGCTGATGGACCA GCACCAAGGACGTTTCTTCAGAAGAAAAGCTGTCTTTCCCCTGTTCTTTACATTCTTTTGCATAAGGCAGGTA GGAACCAAGACAGCTGGGTGGGAAGGGCTCCCTGTCAAAACTCCAACCAGCCTGTGCACTGGGAGGAATGAAC CCTGAGATTCAAACTACGAGTTTACAAATTTGAGGAAGAGCTTTGGTCCAGGGCAGGCTTGGGGAAGAAGTG ACAACCACTCATCTAGGCAGATGCCCTGGGGTGCCGCTGGGGTGGCATGCCAGCATCCATGTAAACTGCCAG CCAAGGACAAATTCATTCCCCAAGGACAAATTTGTCCCCAAGGACAAATTGAAGGTGATATTGTCCCTGCTGACA TCCCTGTCTCAGTTTTTCCTCAATGATTCTCAACCAGAGGAAGCAATACCTCCTCAATCCCTGCTCCGGGTTCC <u> AATTGTTGCAGAGTTGACACCTCCAAAATTGTCATTTGGTTTCCTGAACACAGTTCAGAGTTCAGTACTTCCTACT</u> ATGTATGAACTAGACCGATTATTI

GTCATCAGGACTAAGTGGTGGAGCATCAAAAGGTAGAAAGATGGAACTTATTCAGCCAAAGGAGCCAACTTCACA TACATTAAGTGCCATCAAAAGTCAAAATTATCACTGCACAACTCGTAGTGGGAGGTTTTTACGGTCCATGGGTACA AAGAAGAACCATTTCAGGAGAAAATAGTAGTAAAAAGAGAAGTAAAGAAAAGACGGCAGGAATTGGAGTATCAAG TAAAGCTTGATCTGAACCAATGGCTGGCTATGACTGCTAGCGAGAGACAAGATCTTGTAAGCATCCACCAGTCA GTACATTICICTTTGTCATGAATTGCATACTTTGTTCCAAGTCATGTGGTCTGGAAAGTGGGCGTTGGTCTCACCA **GGCACAGGAGAAGGATACTAATGGGTAAAATCTTTCGAACATGGTTTGAACAATCACCCATTGGAAGAAAAAGC** GAAATGGCACTGTGTGGACTGCAACACGACCGAGTCCATTTGGGCTTGCCTTAGCTGCTCCCATGTTGCCTGTG GAAGATATATTGAAGAGCATGCACTCAAGCACTTTCAAGAAAGCAGTCATCCTGTTGCATTGGAGGTGAATGAGA TGTACGTTTTTTGTTACCTTTGTGATGATTATGTTCTGAATGATAACGCAACTGGAGACCTGAAGTTACTACGACG AAATAGTTTCTGTTCAGGTGCCAGCACAAACGCCAGCATCACCAGCAAAAGATAAAAGTACTCTACCTCAGAAA GAAATTTGGGAAATACTTGCTATATGAATTCTGTTCTTCAGGTGTTGAGTCATTTACTTATTTTCGACAATGTTTT ATGAAATATCTCAAAAAGTCAGTGACTCCTCAGTTAAACGAAGGCCAATAGTAACTCCTGGTGTAACAGGATTGA **GGTGATGATTCTTATTTCTTACATGACGGTGCCCAATCTCTGCTTCAAAGTGAAGATCAACTGTATACTGCTCTTT** ATGCTAGCAATGGATACGTGCAAACATGTTGGGCAGCTGCAGCTTGCTCAAGACCATTCCAGCCTCAACCCTCA TTGCTATGCTACACTCAGTGTGGAGACTCATTCCTGCCTTTCGTGGTTACGCCCAACAAGACGCTCAGGAATTT CTTTGTGAACTTTTAGATAAAATACAACGTGAATTAGAGACAACTGGTACCAGTTTACCAGCTCTTATCCCCACTT >SGPR453 SEQID 10

CTCAAAGGAAACTCATCAAACAAGTTCTGAATGTTGTAAATAACATTTTTCATGGACAACTTCTTAGTCAGGTTACA GATTCCAAACTAAGCATGTGCACTATGGATGAAGTATGCAAGGCTCAAGCTTATATCTTGTTTATACCCAACGAG TGTCTTGCATGTGACAACAAATCAAATACCATAGAACCTTTCTGGGACTTGTCATTGGAGTTTCCAGAAAGGTATC TTACTGAGAATGGACATTCTAAACTTTTGCCTCCAGAGCTCCTGTTGGGGAGCCAACATCCCAATGAAGACGCTG CTTTAGAAGGAAAAATCTACGTATGTGACCAGTGTAACTCAAAGCGTAGAAGGTTTTCCTCCAAACCAGTTGTACT GTCAGGACGTAATAACCGAGAGAAGATTGGTGTTCATGTTGGCTTTGAGGAAATCTTAAACATGGAGCCCTATTG AATGCAGTGGAAAAGATATTGCTTCCCAGCCATGTCTGGTTACTGAAATGTTGGCCAAATTTACAGAAACTGAAG CTGCAGGGAGACCCTGAAATCCCTCAGACCAGAATGCTTTATCTATGACTTGTCCGCGGTGGTGATGCACCATG GGAAAGGATTTGGCTCAGGGCACTACACTGCCTACTGCTATAATTCTGAAGGAGGGTTCTGGGTACACTGCAAT ATACCTCGTCTAATGAAATCCTTAGCTGA

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AAATACAGAAGGGAGGAAAATGCAGAAATTTATCTGTAAGAGGAATTACAAATTTAGGAAATACTTGCTTTTTAAT GAAGACTCTTCAGATGATATTGCTGTAGGTTTAACTTGCCAACATGTAAGTCATGCTATCAGCGTGAATCATGTAA TCAAGATTTTTCCTTCCTCAGACTCTCAGCTGGACCCATTGGTGGTGGAACTTTCAAGGCCTGGACCACTGACCT <u> ATGCGGGTGAAAGATCCAACTAAAGCTTTACCTGAGAAAGCCAAAAGAAGTAAAAGGCCTACTGTACCTCATGAT</u> AAACATGCTTCTAAAACACAAACAAGTGCATTTTCTAGAATCATGAAACTTTGTGAAGAAAAATGTGAAACAGATG CAGCCTTGTTCCTGTTTCTTCACAGCATGAAGGAGTGAAAAAAGGACCACTTTCTCCTAAAGGTTCTTTTAATCA TAGTACTTACTTCTGATATTTGGTTGTGCCTCAAGTGTGGCTTCCAGGGATGTGGTAAAAACTCAGAAAGCCAAC GTGTTATGAATGTGAAAAAATTATCAACGCATTGTAATAAGAAGGTTTTGGCTCAGATAGTTGATTTTCTCCAG GCTTTGTCAGAAGCGGGTGCATCTACATTTAATATAA

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<u>ATGACTGTCCGAAACATCGCCTCCATCTGTAATATGGGCACCAATGCCTCTGCTCTGGAAAAAAGACATTGGTCCA</u> GAGCAGTTTCCAATCAATGAACACTATTTCGGATTGGTCAATTTTGGAAACACATGCTACTGTAACTCCGTGCTTC TTGCTGACGTGCCTGGCGGACCTTTTCCACAGCATTGCCACACAGAAGAAGAAGGTTGGCGTCATCCCACCAAA GAAGTTCATTTCAAGGCTGAGAAAAGAGAATGATCTCTTTGATAACTACATGCAGCAGGATGCTCATGAATTTTTA

ATGGCAACATGAACGAACCTGCGGAAAATAATAAACCAGAACTCACCTGGGTCCATGAGATTTTTCAGGGAACGC TGTGGAGCAGAATACATCCATTACCCACTGTCTAAGAGACTTCAGCAACACAGAAACACTGTGTAGTGAACAAAA GGCCCTGCACCTAAAGCGGTTCAAGTACATGGAGCAGCTGCGCAGATACACCAAGCTGTCTTACCGTGTGTCT TCCCTCTGGAACTCCGGCTCTTCAACACCTCCAGTGATGCAGTGAACCTGGACCGCATGTATGACTTGGTTGCG TGTTTGATGATGACATTGTAGAGAAAATAGATGCTCAAGCTATTGAAGAATTCTATGGCCTGACGTCAGATATAT TACCAATGAAACTCGATGCTTGAACTGTGAAACTGTTAGTAGCAAAGATGAAGATTTTCTTGACCTTTCTGTTGA ATATTATTGTGAAACATGCTGCAGCAAACAAGAAGCCCAGAAAAGGATGAGGGTAAAAAAAGCTGCCCATGGTCTT GTGGTCGTTCACTGTGGCAGTGGTCCTAATCGTGGGCATTATATCACTATTGTGAAAAGTCACGGCTTCTGGCTT CAAAAAATTCAGAATCTGGATATATTTTATTCTATCAGTCAAGAGAGTAA

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TAGATATGAGTACTTTTATTGATGTTGAAGATGAGAATCTCCTCAGACTGAAAGTTGCACTGACAGTGGAGCAGA TGGTTATGAGGGCTGGCGAATCGACACATATCTTGATATCCCATTGGTCATCCGACCTTATGGGTCCAGCCAAGC GGTACAGCTTCGATGATCAACATGTCAGCAGGATAACACAAGAGGACATTAAGAAAACACATGGTGGATCTTCAG GAAGCAGAGGATATTATTCTAGTGCTTTCGCAAGTTCCACAAATGCATATATGCTGATCTATAGACTGAAGGATCC TAGTGAGGCTTGGCAGCAGCATGATGTACAAGAACTATGCAGAGTCATGTTTGATGCTTTGGAACAGAAATGGAA GCAAACAGAACAGGCTGATCTTATAAATGAGCTATATCAAGGCAAGCTGAAGGACTACGTGAGATGTCTGGAATG TCTTGAAACCAATAGTGGAACTGAAAAGATCTCAAAATCTGGACTTGAAAAGAATTCCTTGATCTATGAACTTTTCT AATGCATTATATAAGTGGGAATTTGAAGAATCTGAAGAAGATCCAGTGACAAGTATTCCATACCAACTTCAAAGGC CTGAAAAGATTCGATTTTGATTATACAACCATGCATAGGATTAAACTGAATGATCGAATGACATTTCCCGAGGAAC <u> ATGGTGCCCGGCGAGGAGCAACTGGTCCCGAAAGAGGCACCACTGGATCATACCAGTGACAAGTCACTTCT</u> GCTGGAGGATTCCAGTGCTGGGGAAGACAGTGTTCATGACAGGTTTATAGGTCCGCTTCCAAGAGAAGGTTCTG <u> ATTTGCTAGTGTGGAAGAAGCATTGCATGCATTTATTCAGCCAGAGATTCTGGATGGCCCAAATCAGTATTTTTGT</u> GAACGTTGTAAGAAGAAGTGTGATGCACGGAAGGGCCTTCGGTTTTTGCATTTTCCTTATCTGCTGACCTTACAG AAATGAAGGTAGTTGTCACAGTGATCAGATGAGCAACGATTTCTCCAATGATGATGGTGTTGATGAAGGAATCTG GGGACTAGTAAACCAAGCAATGACTTGCTATTTGAATAGCCTTTTGCAAACACTTTTTATGACTCCTGAATTTAGG CTGTTATGGCTCATTCTGGGAGCGCTGCTGGTGGTCATTATTATGCATGTATAAAGTCATTCAGTGATGAGCAGT CGACGCTAATTTTGAGCCAGGAAAGAAGAACTTTCTGCATTTGACAGATAAAGATGGTGAACAACCTCAAATACT TGGGTTCTACCAGTGATTATGTCAGCCAAAGCTACTCCTACTCATCTATTTGAATAAATCAGAAACTGGATATGT

FIG. 1N

AGGTTGTATTGGAAAGCAGTAGTGTGGACGAATTGCGAGAGGAAGCTTAGTGAAATCAGTGGGATTCCTTTGGATG CTGTGGAAGCTATTCTAGAAGAAAGCACTGAAAAACTCAAAAGCTTGTCACTGCAGCAACAGCAGGATGGAGATA CAGCATCAGTGGATAATAGAGAACTTGAACAGCATATTCAGACTTCTGATCCAGAAAATTTTCAGTCTGAAGAACG TTAGAGCCCTTTGTTGGAGTTTTGTCCTCTCACTTCAAGGTCTTTCGAGTGTATGCCAGCAATCAAGAGTTTGAG GTTGGAAGAACAAGAAAAGAGACAACGAGAAATTGAGCGCAATACATGCAAGATAAAATTATTCTGTTTGCATCCT ACAAAACAAGTAATGATGGAAAATAAATTGGAGGTTCATAAGGATAAGACATTAAAGGAAGCAGTAGAAATGGCTI CTAGAACGGTCATATGAAGGAGAAGAAGATACACCAATGGGGCTTCTACTAGGTGGCGTCAAGTCAACATATAT ATCAGACTCAGATGTGAATAATGACAGGAGTACAAGTTCAGTGGACAGTGATATTCTTAGCTCCAGTCATAGCAG TGATACTTTGTGCAATGCAGACAATGCTCAGATCCCTTTGGCTAATGGACTTGACTCTCACAGTATCACAAGTAGT ATATGATGAGAGTGGCAAGAGTAGGGGAGAAATGCAGTACATGTATTTCAAAGCTGAACCTTATGCTGCAGATGA TTAGAGCTCAGTATTGACAGGTTTCGTCTAAGGAAAAAAACATGGAAGAATCCTGGCACTGTCTTTTTGGATTATC ATAAGATGATGGATTTAGAAGAGGTAATACCCCTGGATTGCTGTCGCCTTGTTAAATATGATGATGAGTTTCATGATTA GTTTGATCTGCTGTTGGAGACGAGAAAGCCTGATCAGGTTTTCCAATCTTATAAACCTGGAGAAGTGATGGTGAA TIGTTGAAAGCTCCGAGACTTTGGATTACCAGATGGCCTTTGCAGACTCTCATTTATGGAAACTCCTGGATCGGC GAAAGCTGGAGGCGATTCTGGTAATGTGGATGATGACTGTGAAAGAGTCAAAGGACCTGTAGGAAGCCTAAAAGT AGAAGAACGAAAGCAAATGAAGGGAAAAAAAAAACATGGGATACAGCAGAAGAAGAAGTCTGGAACTGATAGTGA AGGTTCTGGGGAAGGACATAAATGGTTGATGGTGCATGTTGATAAAAGAATTACTCTGGCAGCTTTCAAACAACA TACAATGATTTGCGTCTTCTCAGTGTCTCCAGTAAAACCCTGAAAGCTGAAGGATTTTTAGAAGTAACAAGGTGT ATGGGGACAGCAGCAAAAGTACTGAGACAAGTGACTTTGAAAACATCGAATCACCTCTCAATGAGAGGGACTCT ATATTTATGAAGAAGATATTAATATTTCCAGCAACTGGGAGGTTTTCCTTGAAGTTCTTGATGGGGTAGAGAAGAT GAAGTCCATGTCACAGCTTGCAGTTTTGTCAAGACGGTGGAAGCCTTCAGAGATGAAGTTGGATCCCTTCCAGG ACAGAATTCAAACAACTGATTTCAAAGGCCATCCATTTACCTGCTGAAACAATGAGAATAGTGCTGGAACGCTGC GAATCCTAAAGTTTCTACCCTGAATGTCTGGCCTCTTTATATCTGTGATGATGGTGCGGTCATATTTTATAGGGAT ATGCAAATACAATCAGATTATTTGTTTTGCTACCTGAACAATCCCCAGTATCTTATTCCAAAAGGACAGCATACCA AGCGTCCGGCTGAATGAGACACTTTCATCATTTTCTGATGACAATAAGATTACAATTAGACTGGGGAGAGACTT ATATTGAATTTGCTAAGGGTAGAGGAACATTTCCCTGTGATATTTCTGTCCTTGATATTCATCAGGATTTAGACTG AAAAAAGGAGAATACAGAGTTAAAGTATACCAGCTTTTGGTCAATGAACAAGAGCCATGCAAGTTTCTGCTAGAT

CAGCTGATTCATGGAAGAGACTATGAAATGGTCCCAGAACCTGTGTGGAGGAGCACTTTATCACTGGTATGGAGCA CAGTTACGTGAATACTACAGAAGAGAGAAATTTTCAGACAACATTTCTACTGCATCTGAAGCCTCAGAAACTGCTGGC CACCTATTCGTCCATCTCTAAGTGAAGGTTTGTTTAATGCTTTTGATGAAAATCGTGACAATCACATAGATTTAAG ATTATTAGAGGATGGTTAGAACGAGAGGAGGTATGGTCTGCAAGCAGGACACACTGGTTTATCATCTCCATG CTTCTCTTCCTGAGACAGCAGCCTGCCACTCGGACACAGCAGTCTAACATCTGGGTGAATATGGGAAATGTACCT CACGAATATCTATCTCAAAGGCTGCGCATTAAAGAGGAAGATATGCGCCTGTGGCTATACAACAGTGAGAACTAC GGACAACCGCACTGATGATATTCCTGAATTACATATGGATCTCTCTGATATTGTAGAAGGCATACTGAATGCACAT GACACCACAAAGATGGGTCATCTTACTCTGGAAGACTATCAGATCTGGAGTGTGAAAAAAATGTTGCCAATGAG CAGTGGTGGCAACAGTGGAAAGAATATGTCAAATACGATGCCAACCCTGTGGTAATTGAGCCATCATCTGTTTTG CTTACTCTTCTGGATGATGAGGATCATAAATTGGAATATTTGAAAATCCAGGATGAACAACACCTGGTAATTGAAG GAGATATCCTGTGGGTTATCAGCCTGTTGCAGGGGACCCCTGGCTGAAAGACAAAAATTTTGCTTCAAGGTATTT TTTTGAACCTCCTTTTCCAGGTGTGTCACATAGTTCTGGGGTTAAGACCAGCTACTCCGGAAGAAGAAGAAGACAA AGCGGCTTTCTGTATTCTGCCACACACGGGGCAGATGTTTGCTTTGCTCGACAACACATAACACTTCTGACAATAAC AACCAGTGTTTGCTGGGAGCCAATGGGAATATTTTGTTGCACCTTAACCCTCAGAAAACCAGGGGGCTATTGATAAT CAGCCATTAGTAACTCAAGAACCAGTAAAGGCTACATCATTAACACTAGAAGGAGGACGATTAAAACGAACTCCA CCCACAGAAAAGGGAGCCACAGGTCTAAGCAATCTGGGAAACACATGCTTCATGAACTCAAGCATCCAGTGTG GATGTTGACCGTGATGGAGTTCTCTCCAGGGTTGAACTGAGAGACATGGTGGTTGCACTTTTAGAAGTCTGGAA **AATGGAGGAAAATACTCATTTGGAACTGCAGCCCATCCTATGGAGCAGGTCGAAGATAGAATTGGAAGCAGCCT** ICTCCGAATGCACCTTTAAAGCGGGTATTAGCCTATACAGGCTGTTTTAGTCGAATGCAGACCATCAAGGAAATT

7AAAACGACTGAAGGATGCTTTCAAGAGGACCTGTGGACTCTCATATTACATGGGCCAGCACTGCTTCATCCGGG

AAGTECTTGGGGATGGAGTGCCTCCAAAGGTTGCTGAGGTGATTTACTGTTCTTTTGGTGGAACATCCAAAGGG

<u> ATGGGTGCCAAGGAGTCACGGATCGGATTCCTCAGCTACGAGGGGGCGCTGAGGAGAGTTACAGATGTAGAGC</u>

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AGTGATACTCCTACTTTCTACCAAACTCTGGCTGGAGTCACACTTTGGAGGAATCAGACATCATTGATCTTGAGA

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CSGSGSLS CSGSCS

TTGGGTGACTGGCTGGAGCCAGGTTAAGCAGCGCTTTTCAGGCTCCACAGCCCAACTCCATGCTGACCCCAGAGC ATCATTATGCACCCCAAGTACTGGGGCCGGGCCTTCATCATGGGTGACGTTGCCCTTGTCCACCTTCAAACACCT GTCACCTTCAGTGAGTACGTGCAGCCCATCTGCCTCCCGGAGCCCAATTTCAACCTGAAGGTTGGGACGCAGTG GATAACTGGAGGATTGCCGGCTCCAGACAGGAAGTGGCCCTGGCAGGTGAGCCTGCAGACCAGCAACAGACAC TGGTGGTCGGAGGATTTGGAAATGACCCGCCATTGGCCCTGGGAGGTGAGCCTCCGGATGGAAAATGAGCACG TGTGTGGAGGGGCCCTCATTGACCCCAGCTGGGTGGTGACTGCGGCCCACTGCAGCCAAGGCACCAAAGAGTA AGCCTTCTCAGGTCCCTGTGCCTCTGCCTGCTCCTGTTCCTGTGCTGGCCGCTGCAGCCCCAGATGGGCTCCT CCCAGTCCAGGGGGAACGCAGTGCGCCCCGCTTCGGTGTTACTTCCCTCAGCCTGTGGCCAGCGGACTTCAAG ATCTGCGGAGGCTCCCTTATCGCCAGACACTGGGTTATAAAGAGGACACCAAACCAATCCAGTTTGTGGTGAGCCC CTCAGTGGTGCTTGGCACCTCCAAGCTGCAGCCCATGAACTTCAGCAGGGCCCTCTGGGTCCCTGTGAGGGAC TGCAGGAGGCTGAGGTGTTTATCATGGACAACAAGAGGTGTGACCGGCATTACAAGAAGTCCTTCTTCCCCCCA AAGGCACAGAATCCAGGTGTGTACACCCGCATCACCAAATACACCAAATGGATCAAGAAGCAAATGAGCAATGG GTTGTCCCCCTTGTCCTGGGGGACATGATCTGTGCCACCAATTATGGGGAAAACTTGTGCTATGGGGATTCTGG

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ATGGGCGCGCGCGCGCTGCTGCTGCTGCTGCTGCTCGGGCTGGACTCGGGAAGCCGGAGGCCTG >SGPR447 SEQID 26

TCCGGAGGGTGATGTCCCACAGTACACCCAGGCCAAACCCCTCCCAGCTGTTGCTGCTCCTTGCCTGCTGTGG GGCACACCTCTGCCACCTCCTTACAACCTCCGGGAAGCACAGGTCACCATCTTAAACAACACCCGGGTGTAATTAC CTGTTTGAACAGCCCTCTAGCCGTAGTATGATCTGGGATTCCATGTTTTGTGCTGGTGCTGAGGATGGCAGTGTA GAGCTGGGGAATGGACTGCGGTCAACCCAATCGGCCTGGTGTCTACACCAACATCAGTGTGTACTTCCACTGGA CGCACTGCTTCCAAAAGCACTACTATCCCTCCGAGTGGACGGTCCAGCTGGGCGAGCTGACTTCCAGGCCAACT CCTTGGAACCTGCGGGCCTACAGCAGTCGTTACAAAGTGCAGGACATCATTGTGAACCCTGACGCACTTGGGGT GACACCTGCAAAGGTGACTCAGGTGGACCCTTGGTCTGTGACAAGGATGGACTGTGGTATCAGGTTGGAATCGT TTACGCAATGACATTGCCCTGCTGAGACTGGCCTCTTCTGTCACCTACAATGCGTACATCCAGCCCATTTGCAT CGAGTCTTCCACCTTCAACTTCGTGCACCGGCCGGACTGCTGGGTGACCGGCTGGGGGGTTAATCAGCCCCAGT

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CCGGCGGTCTTCACCGCGCTCCCTGCCTATGAGGACTGGGTCAGCAGTTTGGACTGGCAGGTCTACTTCGCCG CCGGGGGATGGTGTACCAGTGCTGTGGGTGAGCTGCCCAGCTGTGAGGGGCCTGTCTGGGGGCACCACTGGTG CATGAGGTGAGGGGCACATGGTTCCTGGCCGGGCTGCACAGCTTCGGAGATGCTTGCCAAGGCCCCGCCAGG

OTENTATE OTENTAL

>SGPR529 SEQID 28

GGGATGACCACCTGCTTCTTCAGGGCGAGCAGCTCCGCCGGACGACTCGCTCTGTTGTCCATCCCAAGTAC AGGTTGCTGGCTGGGCCCCCCGCCCCGGAGAGTGAAGTACAACAAGGGCCTGACCTGCTCCAGCATCAC <u>ATGAGAGČTCCGČACCTCCACCTCTCCGCCGCCTCTGGCGCCCGGGCTCTGGCGAAGCTGCTGCCGCTGCTGCTGA</u> CACCAGGGCTCAGGCCCCATCCTGCCAAGGCGAACGGATGAGCACGATCTCATGTTGCTAAAGCTGGCCAGGC ACCGGGGCCAGGACCCTTGCCAGAGTGACTCTGGAGGCCCCCTGGTCTGTGACGAGACCCTCCAAGGCATCCT CCGTAGTGCCGGGGCCCCGGGGCCCTGCAGCTTCCCTACCGCTGTGCTCAGCCCGGAGACCAGTGCC GCGCCCCGTGCGCGCGCGCGCCAGCCCTGGCAGGTCTCGCTCTTCAACGGCCTCTCGTTCCACTGCGCGG TATCCTGAGCCCTAAAGAGTGTGAGGTCTTCTACCCTGGCGTGGTCACCAACACATGATATGTGCTGGACTGG TGGCGCAACTCTGGGCCGCAGAGGCGGCGCTGCTCCCCCAAAACGACACGCGCTTGGACCCCCGAAGCCTATG GTGTCCTGGTGGACCAGAGTTGGGTGCTGACGGCCGCGCACTGCGGAAACAAGCCACTGTGGGGCTCGAGTAG CTCGTGGGGTGTTTACCCCTGTGGCTCTGCCCAGCATCCAGCTGTCTACACCCCAGATCTGCAAATACATGTCCT GGATCAATAAAGTCATACGCTCCAACTGA

>SGPR428 1 SEQID 29

GGTCATGGGAACGAGAACATTCAGCAACATCCACTCGGAGAGAAAGCAAGTGCAGAAGGTCATTATTCACAAATA AATGCCTGTCTGCCTGCAGGAGGAGGAGGACCTGGGACTGGTGTTGGATGGCACAGTGGGTAACGACCAAT CTGGGGTGTGGGCCTGTGGCCAGAAGAACATGCCTGGAGTGTACACCGAGTTGTCCAATTATCTGCTTTGGATCG GGGTATGACCAATATGATGACTTAAACATGCACCTGGAAAAGCTGAGAGGTGGTGCAGATTAGCCGGAAAGAATG AAAACGTACAATGTGGCCACAGGCCTGCTTTTCCAAACTCGTCATGGTTACCATTTCATGAACGGCTTCAAGTCC AGAATGGTGAGTGCCCGTGGCAAGTGAGTATCCAGATGTCACGGAAACACCTCTGTGGAGGCTCAATCTTACAT TGGTGGTGGGTTCTGACAGCCGCACACTGCTTCCGAAGAACCCTATTAGACATGGCCGTGGTAAATGTCACTGT TTACAAACCGCCCCAGCTCGACAGTGACCTCTCTCTGCTTCTACTTGCCACACCAGTGCAATTCAGCAATTTCAA

AGAGGAAGACTGTGCTGGCAGGGAAGCCGTATAAGTATGAGCCAGACTCTGTGTACGCTTTGCTTCTCTCACCC TGGGCCATCCTGTTACTGTATTTTGTGATGCTTCTATTATCCT

>SGPR425 SEQID 30

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FIG. 1DD

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TGTTGCTGCGCCTAGTCCAGCCCGCACGCCTGAACCCCCAGGTGCGCCCCGCGGTGCTACCCACGCGTTGCCC CCACCCGGGGGGGGCCTGTGTGTGTTCTGGCTGGGCCTGGTGTCCCACAACGAGCCTGGGACCGCTGGGAG ATGGGGGACCCAGAAGGAAGCGCAGAGTGGGGTTGGGGGAAGGGGGATACCGGTGGTCAGAAGAAATTTATTAA CAGTGGATGGGATAAGTCTGTGTCTGGAGGGATCCTGGTGGAGGCAGAAGGGTCCTGCCTCACCTGGATTCTCT GGCAAGTGGCTCTCTACGAGCGTGGACGCTTTAACTGTGGCGCTTCCCTCATCTCCCCACACTGGGTGCTGTCT GAGCAACTACGGACCACGTCTCGGGTCATTCCACACCCGCGCTACGAAGCGCGCAGCCACCGCAACGACGTCA CTGGCATCCACAGCAGCCCAGGATGGTGACAAGTTGCTGGAAGGTGACGAGTGTGCACCCCCACTCCCAGCCAT GCGGCCCACTGCCAAAGCCGCTTCATGAGAGTGCGCCTGGGAGAGCACAACCTGCGCAAGCGCGATGGCCCA

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>SGPR396 SEQID 32

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>SGPR426 SEQID 33

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>SGPR485 1 SEQID 36

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ATCCAGGTCCTGAAGATCGCCAAGGTCTTCAAGAACCCCAAGTTCAGCATTCTGACCGTGAACAATGACATCACC CAAGCTGCAGCAGCCAGCCCTGCCCTCCTGTCCAATGCCGAATGCAAGAAGTCCTGGGGCAGGAGGATCACC GACGTGATGATCTGTGCCGGGGCCAGTGGCGTCTCCTCCTGCATGGGTGACTCTGGAGGCCCCCTGGTCTGCC ACTTCCCCGCGGGGACACTGTGTGCCACCACAGGCTGGGGCAAGACCAAGTACAACGCCAACAAGACCCTGA GCCCACTGCGGGGTCAGGACCTCCGACGTGGTCGTGGCTGGGGAGTTTGACCAGGGCTCTGACGAGGAGAAC CCCTGTGCTCAGCGGCCTGTCCAGGATCGTGAATGGGGAGGACGCCGTCCCCGGCTCCTGGCCTGGCAGGT GTCCCTGCAGGACAAAACCGGCTTCCACTTCTGCGGGGGCTCCCTCATCAGCGAGGACTGGGTGGTCACCGCT AGAAGGACGGAGCCTGGACCCTGGTGGGCATTGTGTCCTGGGGCCGCCGCCCTGCTCTACCACCACGCCCG CTGTGTACGCCCGTGTCACCAAGCTCATACCCTGGGTGCAGAAGATCCTGGCCGCCAACTGA >SGPR534 SEQID 37

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>SGPR390 SEQID 38

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FIG. 1KK

GCCTGGGTCCTGGTGGGGCTGGCCAGCTGGGCCCTGGACTGCCGGCATCCTGCCTACCCCAGCATCTTCACCA GAAGACCATAAGAGGGTGCAACTGTCACCACCCTTCTATCTCCAGGAGGGCAAGGTGGGCCTCATTGAGAACAC GGGTCACCTACTTCATCAACTGGATTGACGAAATCATGAGGCTCACTCCTTTCTGACCCCGCGCGTGGCTCCTC GGTATCAACTACCCACTGCTTTCTCAACAAATCCCAGGCCCCGAAGAACTATCAGGTTCTGTTGGGAAACATCCA GGGGGTACTTCTCGACAGGAAAGTCCATCTGCAAAGGCGATTCTGGGGGGCCTCTAGTCTGCTACCTCCCCAGT >SGPR530_1_SEQID_40 GTCTCCACAGTGTGTGGGAAGCCTAAGGTGGTGGGGAAGATCTATGGTGGCCGGGACGCAGCAGCTGGCCAGT ACTGTATCATCAAACCCAGCACACCCAGAAGATGTCTGTGCACCGGATCATCACCCATCCAGACTTTGAGAAGCT ACTCTGTAATACCTTATATGGGCAAAGAACTGCAAAGGCGAGACCTAAGCTTTGCACGAGGAGATGCTGTGTGG CCACCCCTTTGGGAGTGACATTGCCATGTTGCAGCTGCACCTGCCTATGAACTTCACTTCCTACATTGTCCTGT CTGCCTCCCATCCCGGGACATGCAGCTGCCCAGTAACGTGTCCTGTTGGATAACCGGCTGGGGAATGCTCACC GGCCATGGCAGGCCAGCCTGCTCTACTGGGGCTCGCACCTCTGTGGAGCTGTCCTCATCGACTCCTGCTGGCT

>SGPR520 SEQID 41

ATGCTGCTGGCTGCTGCTGCTGCTACCCCTCCCAAGCTCATGGTTTGCCCACGGGCACCCACTGTACACACG CCTGCCCCCCAGCGCCCTGCAAGTCTTCACTCTCCTTGGGGGCAGAGACTGTGTTGGGCCGCAACCTAGACT ACGTTTGTGAAGGGCCGTGCGGCGAGAGGCGTCCGAGCACTGCCAATGTGACGCGGGCCCACGGCCGCATCG TGGGGGGCAGCGCGCGCCCCGGGGCCTGGCCCTGGCTGAGGCTGCAGCTCGGCGGCGGGCAGCTCTG CCTGCCGGAACCGCCTGCGCCATCGCGGGCTGGGGCCCCCTCTTCGAAGACGGGCCTGAGGCTGAAGAGGTG SCCCGGCTCTGCGCCTTCTATGCCCGCCTGTGCCCGGGGTCCCAGGGCGCCTGTGCGCGCCTTGTGCGCGCCTGGCGCCCAG AGAGAGGCCCGTGTTCCCCTGCTCAGCACCGACACCTGCCGAAGAGCCCTGGGGCCCGGGCTGCGCCCCAGC ACCATECTCTGCGCCGGGTACCTGGCGGGGGGGGGTTGACTCGTGCCAGGGTGACTCGGGAGGCCCCTGACC TGCGGCGCGCCTCCTGGTAGCGGCCTCCTGGGTGCTCACGGCAGCGCACTGCTTTGTAGGCTGCCGCTCGACC GTGCCAGTGAACCGCATCCTGCCCCCACCCCAAGTTTGACCCGCGGACCTTCCACAACGACCTGGCCTGGTGC TGTTCTGAGCCTGGCCCCCGCCCTAGAGAGGTCCTGTTCGGAGTCACCTCCTGGGGGGGCGGCTGCGGGGGAG

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TGTGGAATGTTGACCATCACTTCAATATTCAGTTCTAGTAACATGACGGTGATATACTTTAAAAGTGATGGTAAAA AATATTCAAATGTCCTACAAGAAATGGAACTTCCCATCATGGATGACAGAGCGTGTAATACTGTGCTCAAGAGCAT AAATTTAGTCCAAACTGTATTTATGATGCTGTTGTGATTTACGGTGATTCTGAAGAAAAGCACAAGTTAGCTAAACT GTGAGAAGGGCCAAACACATAATAGTGCATGAAGACTTTAACACACTAAGTTATGACTCTGACATTGCCCTAATAC ATTTACAAGTTTAGACATGGAAAAGCAAGTTGGATGTGATCATGACTATGTATCTTTACGATCAAGCAGTGGAGTG TCCCCAAAACAATCCTGTATCTACCGTAAAAGCTATTCTGCATGATGTCTGTGGCATCCCTCCATTTAGTCCCAG AGGTTGTGGGAGTCTGGCTATATTGGTAGAAGAAGGGACAAATCACTCTGCCAAGTATCCTGATTTGTATCCCAG ATGAGTCCTGATATTGCACTGCTGTATCTAAAACACAAAGTCAAGTTTGGAAATGCTGTTCAGCCAATCTGTCTTC CTGACAGCGATGATAAAGTTGAACCAGGAATTCTTTGCTTATCCAGTGGATGGGGCAAGATTTCCAAAACATCAG GAACCTCCCTCCCCTGGGAAGGACCATGCTGTGTGCTGGCTTCCCTGATTGGGGAATGGACGCCTGCCAGGGG GACTCTGGAGGACCACTGGTTTGTAGAAGAGGTGGTGGAATCTGGATTCTTGCTGGGATAACTTCCTGGGTAGC TGGTTGTGCTGGAGGTTCAGTTCCCGTAAGAACAACCATGTGAAGGCATCACTTGGCATTTTCTCCAAAGTGTC TGAGTTGATGGATTTTATCACTCAAAACCTGTTCACAGGTTTGGATCGGGGCCAACCCCTCTCAAAAGTGGGCTTC TAACACAAGGTGTCATTGGTTCATTTGTGCTCCAGAGAAGCACATTATAAAGTTGACATTTGAGGACTTTGCTGTC TGAAGAATAATCCACTCTCCTGGACTATTATTGCTGGGGACCATGACAGAAACCTGAAGGAATCAACAGAGCAG AAGGTATATAACAAAGGCCCTGAGTTCTGTCCAAGAAGTGAATGGAAGCCAGAGAGATAAAATAATCCTGATAAA ATCGTTTACAAGGCTTCAAGGCCAGATTTACCATTTTGCCCTCAGAGTCTTTAAACAAATTTGAACCAAAGTTACC GGCTTTCCAGAAGAATCGCAGGAGGGGAAGAAGCCTGCCCCCCATGGTTGGCCATGGCAGGTGGGTCTGAGGT TCTAGGCGATTACCAATGTGGAGGTGCCATCATCAACCCAGTGTGGATTCTGACCGCAGCCCACTGTGTGCAA GTTTCTGATACAGAGACAGTGGCAGTGGCTTTGAGCTTACCGTTACTGCTGTACAGAAGTCAGAAGCAGGGTC CTITITAGTAAGGTCTGTGGAAAAAATATTGCCTTCACCATTGCTGGCAGAGACCAGTGAGGCCATGGTTCCATTT

FIG. 1LL

ATGCAGATCCCAAGAAATTCTCACAGAACTACTATGGGTTGCCAATGGAGATTAGTAGCCCCTTTAAATCACATCA AGACTTTCCCAAAGTTCAAACAGAGAGCACTTGGTCCCTTGTGAGGATGTTCTTCTGACCAAGCCAGAAGGGATC ATTTGGACCAGGAAAAAAATTAATAGGTAGAATGTTGATGAGCACTGAGCTTTCTTGGTTCCTAAGCCAATTCAGC GGTGGATTACATGGAAGAAGACTTTACTCAATGACTTTCATGAGTCCTGGACCGCTGGTGAGGGTGACATTCCAT ACCAAGAAGACCACAGCTTCTTGTGGGGAGACTGCAGTATCTATGAAAATGATGTATACTTCTATCTTGTTGTTGCCC CTGCCCATCCAGGGGGATCACAGAGAAGATGATCTGTGCTGGCTTTGCAGCATCTGGAGAGAAAGATTTCTGC GGGAGCTGGCTGTCCAGCCATGGAAGCCGGGTGTATTTGCCAGAGTGATGATCTTCTTGGACTGGATCCAAT AGTTGTTGCTCTGAAGCAGAGGCTAGAAAGCCTAGAGGCTTTTTTCCCACACCACGGTATCTACTGGATTATAGA GCCCTTGTACGAGGTGCATTTGGTATAAGCTATATTGTCTTGAAAGTCCTAGGTCCAAAGGACAGTAAAATAACC CAGGGAGACTCTGGTGGGCCACTAGTATGTAGACATGAAAATGGTCCCTTTGTCCTCTATGGCATTGTCAGCTG GTGGCCTAGCAAGTCGCCTACAGCAGATTCAAGTGCATGTTAGAAAGAGAGGGTCTGTGAACACACTTACTAT TACAGAACACCTGTTACCATGCACTGCCTCATGAGGTTGTTTTGAGAATTAAATAA

TGGCCCCAGCTCACTGCTATTTACCAAATCTGAAAGTGATGCTGGGAAATTTCAAGAGCAGAGTCAGAGACGGTA TGCTCATCAAGCTGGCTAAGCCTGCCATGCTCAATCCCAAAGTCCAGCCCCTTACCCTCGCCACCACCAATGTCA GGCCAGGCACTGTCTGTCTACTCTCAGGTTTGGACTGGAGCCAAGAAAAAAGAGTGGGCTTTGGCAGCTGGAGCCA CTGCTCCCTATTTGGTGTACCTCAAGTCTCACTTCAACCCCTGTGTGGGCGTCCTCATCAAACCCAGCTGGGTGC CTGAACAGACAATTAACCCCATTCAGATCGTCCGCTACTGGAACTACAGTCATAGCGCCCCACAGGATGACCTCA CCAGGCCATCTGACTCTGCACAGAGGCCCAGCCATTCCTGATTGGCAGAGACACAATTCACATGAACAAGGCCG AAAGCCACAGGAATTCCTTATGTGAAATTTGTGAAAGTATTCAGCCGAATTTTTGGGGAGGTGGCCGTTGCTA >SGPR507_2_SEQID_43

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GTAGCAGCAACTGGAATGACTCCTACTCAGAGAGACCTGCCAGCAGCTGGGTTTCGAGAGTGCTCACCGGACA CTCCAGCTGGGACACCTCCAGGCCGGGCATCTCCAGCCCAGGCATCTCCAGCCCAGGCATCTCCAGCTGGGAC CAGAGTGTACCTTGTTAGAGCAACACCAGTGGGGGCTGTACCCATCCGATCATCTCCTGCCAGGTCAGCACCAG CTGGCATCACTTTCCAGGTCCTCATCCGGCAGGTCATCCTCCGCCAGGTCAGCCTCGGTGACAACCTCCCCAAC GCTGCGTGAGGTTTGACTGGGACAAGTCTCTGCTTAAAATCTACTCTGGGTCCTCCCATCAGTGGCTTCCCATC GGCATCTCCAGCCCAGGCATCTCCAGCCCAGGCATCTCCAGCCCAGGCATCTCCAGCCCGGGCATCTCCGGCT CAACCAGGGCCACCAGGGAGAGCCCAGTCCAGTTCTGGCAGGGCCCACACAGGGATCAGGTACAAGGAGCAGA GGGAGAGCTGTCCCAAGCACGCTGTTCGCTGTGACGGGGTGGTGGACTGCAAGCTGAAGAGTGACGAGCTGG <u>ATGGAGAGGGGACAGCCACGGGAATGCATCTCCAGCAAGAACACCTTCAGCTGGAGCATCTCCAGCCCAGGCAT</u> ACCTCCGGGCCGGGCATCTCCAGCCCAGGCATCTCCAGGTACACCTCCAGGCCGGGCATCTCCAGGCCG >SGPR567 1 SEQID_45

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 IGCTTTAGGAAAAAAAAAAAATGTGATGGGACAGTGGATTGTCCAGATGGAAGTGAAGAAGAAGGCTGCAC
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CGGACAAAGCAACACATATCAACTTAAGGACTTACGAGAGGGCGACGGAAATTTGGTGTATTCTTTGAAAATGTAC ATGACATTGAACAAAATTAAAAGACCTTTTTGCAGGGAAAGGACAGTGGGATTTGGCACCCGAAGCAGAAATGCTG AAGCCATGGATGATTGCCGTTCTCATTGTGTTGTCCCTGACAGTGGTGGCAGTGACCATAGGTCTCCTGGTTCAC CTTTCTTTTGTGTGTCACAGTCCAGAGGAAGATGGTGTGAAAGTAGATGTCATTATGGTGTTCCAGTTCCCCTCTA AAGCAAGTTGTGGTAAACGAGTTGTTCCATTAAACGTCAACAGAATAGCATCTGGAGTCATTGCACCCAAGGCGG AAACCTGGAATGTTCTGTGCCGGATATATGGAAGGAATTTATGATGCCTGCAGGGGGTGATTCTGGGGGACCTTTA GTCACAAGGGATCTGAAAGATACGTGGTATCTCATTGGAATTGTAAGCTGGGGAGATAACTGTGGTCAAAAGGAC TGCATCCTTCCAACCAAATTTGACTGTCCACATCACAGGATTTGGAGCACTTTACTATGGTGGGGAATCCCAAAAT CTGAACAAAGGGCAGTAAGAGAAGAAAAATCCAAAGCATCTTAAATCAGAAGATAAGGAATTTAAGAGCCTTGC GGAGAGGCTCCAGGCCTGGGAGCAGGTCCTGCTGGTCACCAATGAGCTCATCAACAGGGGGAGTTAACTGTCC TTGTCACTGCAGCACACTGCTTCCAGAAGTATAAAAATCCACATCAATGGACTGTTAGTTTTGGAACAAAAATCAA CAATAAATGCCTCATCAGTTCAAGTTAATGTGGCCATGGTCAAGAATGGCAATGTGGGGGCCAGGTTCCGGAGCA CCCTCCCTTAATGAAAAGAAATGTCAGAAGATTTATTATCCATGAGAAGTACCGCTCTGCAGCAAGAGAGTACGA TTCCTAGTATTTGACCAAAAAAAGGAGTACTATCATGGCTCCTTTAAAATTTTAGATCCACAAATCAATAACAATT AAGCCTGGAGTCTACACACAAGTGACTTATTACCGAAACTGGATTGCTTCAAAAACAGGCATCTAA

FIG. 1RR

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GCAGTCTGTGGCTCCTGGGCGCTGGCCGTGGCCAGCGTGGCCCTGGGCTTCCGGCACACGTGTGGGGG CCTGCTGCCTCTCAGCCCATTTCCGGGACCTTGCAGGATGAGGAGATAACTTTGAGCTGCTCAGAGGCCAGCGC CTGCTGGAGCCTTGGGCATCTCAGACTCACTCACCACAGGGAGTAAACCTCACTGACATCAAACTCAACAGTTC CTCTGTGCTAGCGCCACGCTGGGTGGTGACTGCTGCACATTGTATGCACAGTTTCAGGCTGGCCCGCCTGTCCA TGAGGAAGCTCTGCTCCCTGCACTCCCCAAAACAGTATCTTTCAGAATAAACAGCGAAGACTTCTTGCTGGAAGC ACTTCTGGTCAAGTTGTTTCCCTCAGATGCTCTGAGTGTGGAGCGAGGCCCCTGGCTTCCCGGATAGTTGGTGG TCAGAGCAGAGCCTGGAGACCAGCAGCATCCCATTTCTCAGGCGGTGTGCTGGCGTTCCATGCGACGTGGCTG TGCAGTGCTGGGAGCCCTGGGGCTGCTGGCCGGTGCAGGTGTTGGCTCATGGCTCCTAGTGCTGTATCTGTGT CCAGGAGTTTGCTCAGCTCTCTCTAGACTGGGAGGCTTCCTGGAGGAGGCGTGGCAGCCCAGGAACAACTGC GCTGGCGGGTTCATGCGGGGCTGGTCAGCCACAGTGCCGTCAGGCCCCCACCAAGGGGGCTCTGGTGGAGAGGA GGGTGTCTGGCTGGGGCCACCCACCCTAGCCATACTTACAGCTCGGATATGCTCCAGGACACGGTGGTGCC TTATCCCACACCCCCTCTACAGTGCCCAGAATCATGACTACGACGTCGCCCTCCTGAGGCTCCAGACCGCTCTC GCTACCTGGACGGAAGGGCTGATGCATGCCAGGGAGATAGCGGGGGCCCCCTAGTGTGCCCAGATGGGGACA ATGAGCCTGATGCTGGATGACCAACCCCCTATGGAGGCCCAGTATGCAGAGGAGGGCCCAGGACCTGGGATC1 GCAAGTGAGGGATCAGCCACGCTGGCTCCTGGTCTGCCATGAGGGCTGGAGCCCCGCCCTGGGGCTGCAGAT AACTTCTCAGACACTGTGGGCGCTGTGTGCCTGCCGGCCAAGGAACAGCATTTTCCGAAGGGCTCGCGGTGCT

>SGPR527_1_SEQID_52

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GGGTTCAGAGCCTGGGCCTTCCCACCCAGCCCCAGAAGACCCAGGTCAGATCCCCAGGAGCCCAGGGAG GAGAACTGCACCATTGCCCTGCCTGAGTGCGGGAAGGCCCCGCGGGCCAGGGGCCTGGCCCTGGGAGGCCCAG GCTTTCTGGACCCGAACAGCTCCGACAGCCCACCCCGCGACCTCGACGCCTGGCGCGTGCTGCTGCTCGCG 'GTGGACGGAGAAACCGCCCTGGAGTTTTCACTGCTGTGGCTACCTATGAGGCATGGATACGGGAGCAGGTGAT CCGCTGCCCGGAGACCCGCCGCCGCGCGCTCTGCCCTGCCTACCAGGAAAAGGAGGAGGTGGGCAGCTGCTGG ACTCATGGCCCATGGATCAGCCATGTGACTCGGGGAGCCTACCTGGAGGACCAGCTAGCCTGGGATTGGGGCC CTGATGGGGAGGAGACTGAGACACAGACTTGTCCCCCACACAGAGCATGGTGCCTGTGGCCTGCGGCTGGA TCACTCTCCAGATATTGCCAGGGATGCTGTGTGCTGGCTACCCAGAGGGCCGCAGGGACACCCTGCCAGGGTGA GTGATGGTGCCAGGATCCAGACCCTGCCATGGGGCGCTGGTGTCTGAAAGCTGGGTCTTGGCACCTGCCAGCT GCCCTCCCCATCAGCCCTGCCCATCTGTCTCCACCCGGGGGTATCCCCCCGGGGGCCAGCTGCTGGGTGTTG GGCTGGAAAGAACCCCAGGACCGAGTCCCTGTGGCTGCTGTCTCCCATCTTGACACAACGAATCTGTGACTG CCTGCCCCCCAGTGGCTCCCCACACTGGCCCACTGGAGGCAGCAATCTCTGCCCCCCAGAACTGGCCAAGGGC CGCTGCTGCAGCTGCGCCCGTGAACCTGAGCGCGGCTTCGCGGGCCCGTGTGCCTACCCCACCCGGAAC CAGCATCCGGCTGCCCCAGCACCTGGGACTCAGGCCCCCCCTGGCCCTCCTGGAGCTGAGCTCCCGGGTGGA CCTCTATCAGGGCATCCTGCCCCCTGGAACCCTCTGTGTCTGTATGCAGAGGGGGCAGGAGAACAGGTGTGAG ATGACCTCAGCACCGCCCTCCTGTGCCAGATGACGGAAGGGTCCTGGATCCTCGTGGGCATGGCTGTTCAAG CCCGCGCGCGCGGGGGGGGCGCCCTGGTGCAGCACGAGAACGCTTCGTGGGACAACGCCTCGGACCTGG ACTACTTCCTGCCCGGGAGCCGCTGCCGCTGGCCCGCTGGGGCCGCGGGGAACCCGCGCGTTGGCCCAGGC GECTECTCCAGTGGGGGTCCTGTGGCCTGGCTGGCAGAGGTGCATGTGGCTGGTGATCGAGTCTGCACTGG GGAGCCGGGAGCTGTTTGCTGCCATTGGTCCTGAAGAGGCCTGGATCTCCCAGACAGTGGGAGAGGCCAACTT TCGGGATCCCCGCATGCAGTCTACTTCCTGCTCCTGCTGACTCTCCTGATCCAGAGCTGA

ATGGCATCCGTGCGCTTCGGGGGCCAACATCACTGCGGAGGCTTCCTGCTGCGAGCCCGCTGGGTGTCTCGG AAGCCCCCCGCAGGCTCCTGGGGGGCCCAGATCATCGGGGGCCACGAGGTGACCCCCCACTCCAGGCCCTAC CCGCCCACTGCTTCAGCCACAGAGACCTCCGCACTGGCCTGGTGGTGCTGGGCGCCCACGTCCTGAGTACTGC GCCATGGGGCTCGGGTTGAGGGGCTGGGGACGTCCTCTGCTGACTGTGGCCACCGCCCTGATGCTGCCCGTG GGAGCCCACCCAGCAGGTGTTTGGCATCGATGCTCTCACCACACACCCCGGACTACCACCCCATGACCCACGCC >SGPR542 SEQID 53

AGGAGCTGCCGCCTGGACTGATGGAGGCCAAGGTCCGAGTGCTGGACCCGGACGTCTGCAACAGCTCCTGGAA AACGACATCTGCCTGCTGCAGCTGAACGGCTCTGCTGTCCTGGGCCCTGCAGTGGGGCTGCTGAGGCTGCCAG GGGCCACCTGACACTTACCATGCTCTGCACCCGCAGTGGGGACAGCCACAGACGGGGGCTTCTGCTCGGCCGAC CCCAAGACCCCCGACGTGTACACGCAGGTGTCCGCCTTTGTGGCCTGGATCTGGGACGTGGTTCGGCGGGGCA TCCGGAGGGCCCCTGGTGTGCAGGAACCGGGCTCACGGCCTCGTTTCCTTCTCGGGCCTCTGGTGCGGCGAC GTCCCCAGCCCGGCCCCTGCCTGGGACCACCAGGCCCCCAGGAGAAGCCGCCTGA

>SGPR551_SEQID_54

GGCCAGTGGACGATCCAGAACAGGAGGCTGTGTGGCTTGCGCATCCTGCAGCCCTACGCCGAGAGGATCCCCG <u> ATGCCCGTGGCCGAGGCCCCCCAGGTGGCTGGCGGGCAGGGGACGGAGGTGATGGCGAGGAAGCGGAGCC</u> CAAAGAGGACAGCACATGCATCTCACTGCCCAAGGTCTGTGATGGGCAGCCTGATTGTCTCAACGGCAGCGATG GTGCTGCTGCCCTGCTCGTGCTGGCTTCGGCGGGGGTGCTACTCTGGTATTTCCTAGGGTACAAGGCGGAGG GTCCACAGTCAACAGCTCGGCTGCCGTCCCCTACAGGGCCGAGTACGAAGTGGACCCCGAGGGGCCTAGTGATC GGACCTCATGCTCAAACTCCGGCTGGAGTGGACGCTGGCAGAGTGCCGGGACCGACTGGCCATGTATGACGTG TGGACTACGGCTTGGCCCTCTGGTTTGATGCCTATGCACTGAGGAGGCAGAAGTATGATTTGCCGTGCACCCAG TGATGGTCAGCCAGGTGTACTCAGGCAGTCTGCGTGTACTCAATCGCCACTTCTCCCAGGATCTTACCCGCCGG TGGTGGCCACGGCCGGGATCACCATCAACTTCACCTCCCAGATCTCCCTCACCGGGCCCGGTGTGCGGGTGTGC AAGAGCAGTGCCAGGAAGGGGTGCCATGTGGGACATTCACCTTCCAGTGTGAGGACCGGAGCTGCGTGAAGAA GAATCTAGTGCCTTCCGCAGTGAAACCGCCAAAGCCCAGAAGATGCTCAAGGAGCTCATCACCAGCACCCGCCT CTGGAAGCCAGTGTGAAAGACATAGCTGCATTGAATTCCACGCTGGGTTGTTACCGCTACAGCTACGTGGGCCA GGGCCAGGTCCTCCGGCTGAAGGGGCCTGACCACCTGGCCTCCAGCTGCCTGTGGCACCTGCAGGGGCCCCAA CTGGCGTCGGGGGCCATCATGGCGGTCGTCTGGAAGAAGGGCCTGCACAGCTACTACGACCCCTTCGTGCTCT CCGTGCAGCCGGTGGTCTTCCAGGCCTGTGAAGTGAACCTGACGCTGGACAACAGGCTCGACTCCCAGGGCGT CCTCAGCACCCCGTACTTCCCCAGCTACTACTCGCCCCAAACCCACTGCTCCTGGCACCTCACGGTGCCTCTC CCTGTGATGGGGTCAAGGACTGCCCCAACGGCCTGGATGAGAGAACTGCGTTTGCAGAGCCACATTCCAGTG GCCCAACCCGCAGTGTGATGGGCGGCCCGACTGCAGGGACGGCTCGGATGAGGAGCACTGTGACTGTGGCCT AGAGGGGATGTTCAAGGCCTGTGAGGACTCCAAGAGAAAAGCCCGGGGCTACCTCCGCCTGGTGCCCCTGTTT CCAAATCCCCGAGCACCGCCGGCTGATGCTGAGCCCCGAGGTGGTGCAGGCACTGCTGGTGGAGGAGGTGCT CTATGGCTTGTACAACCAGTCGGACCCCTGCCCTGGAGAGTTCCTCTGTTCTGTGAATGGACTCTGTGTCCCTG GCCGGGCCCCTGGAGAAGAGGCTCATCACCTCGGTGTACGGCTGCAGCCGCCAGGAGCCCGTGGTGGAGGTT

CCAGGGCCCCTCCAGCCGCATTGTTGGTGGAGCTGTGTCCTCCGAGGGTGAGTGGCCATGGCAGGCCAGGCCTC TCCAGGAGGACAGCATGGCCTCCACGGTGCTGTGGACCGTGTTCCTGGGCCAAGGTGTGGCAGAACTCGCGCTG GACGTGGCGCTGCTGCAGCTCGACCACCCGGTGGTGCGCTCGGCCGCCGTGCGCCCCGTCTGCCTGCCCGCG ATCAGCAACGCTCTGCAGAAAGTGGATGTGCAGTTGATCCCACAGGACCTGTGCAGCGAGGCCTATCGCTACCA GGTGACGCCACGCATGCTGTGTCCCGCTACCGCAAGGGCAAGGATGCCTGTCAGGGTGACTCAGGTGGT CCGCTGGTGTGCAAGGCACTCAGTGGCCGCTGGTTCCTGGCGGGGGCTGGTCAGCTGGGGCCTGGGGCTGTGGC CAGGTTCGGGGTCGACACATCTGTGGGGGGGCCCTCATCGCTGACCGCTGGGTGATAACAGCTGCCACTGCT GCCTGGAGAGGTGTCCTTCAAGGTGAGCCGCCTGCTGCACCCGTACCACGAAGAGAGACAGCCATGACTAC CGCTCCCACTTCTTCGAGCCCGGCCTGCACTGCTGGATTACGGGCTGGGGCGCCTTGCGCGAGGGGGGCCCC CGGCCTAACTACTTCGGCGTCTACACCCGCATCACAGGTGTGATCAGCTGGATCCAGCAAGTGGTGACCTGA

CAGCCTATTTGCCTACCTTTTGATGTTTTCCAAATCCTGGACGGAAACACAAAGTGTTTTATAAGTGGCTGGGGAA GAACAAAAGAAGAAGGTAACGCTACAAATATTTTACAAGATGCAGAAGTGCATTATATTTCTCGAGAGATGTGTAA <u> AGTTACGGACATGGCTGTGGTCGAAGAGGTTTTCCTGGTGTCTATATTGGGCCATCCTTCTACCAAAAGTGGCTG</u> AGTGAGAGAGAGGTGGGTCCTCACAGCTGCCCACTGCACTAAAGACACTAGCGATCCTTTAATGTGGACAGCTG TGATTGGAACTAATAATACATGGACGCTATCCTCATACCAAGAAGATAAAAATTAAAGCAATCATTATTCATCCA TTCTGAGAGGAGTTATGGGGGAATAATTCCTAACACTTCATTTTGTGCAGGTGATGAAGATGGAGCTTTTGATACT AGGATTGTGGAACAGCACCGCTTAAGGATGTGTTGCAAGGGTCTCGGATTATAGGGGGCACCGAAGCACAAGCT AACTTCATTTTGGAATCTTATGTAAATGATATTGCACTTTTCACTTAAAAAAAGCAGTGAGGTATAATGACTATATI TGCAGGGGTGACAGTGGGGGACCATTAATGTGCTACTTACCAGAATATAAAAGATTTTTTGTAATGGGAATTACC GACCTGCCGCCATCTTGCTCACCAGCCTCCAAAATGCGGCTGGGGCTCCTGAGCGTGGCGCTGTTGTTGTGG GGAGCTCTCACTTATACTCAGACCACTACTCGCCCTCTGGAAGGCACAGGCTCGGCCCCTCGCCGGAACCGGC TATGTTTTGTCATCTTACTAGCAACAACATAA >SGPR451_SEQID_55

TGGGCAGCAGCTTGAAGTGGGCGTGGGCCAAGCCCTCTGGGATGCCCGTCCCAGAGAATGACCTGGTGGGCAT CAGCCCTCAGACTGGCTGTGCTTCGCGGATCTTGGCTGGTTCGAGGCTGATGGAGCTGCCCACTCCATGGGCC AGCCCCCGCAGCCCAGGACCCCTGACTGTAGGCTCCAGGCCTCCCTGGAAGCCCTGGCCACGCTCGCCCCG >SGPR452 1 SEQID 56

GGTGCTGGAGAACGCCGTCTGTGAGCAGCCCTACCGCAACGCCTCAGGGCACACTGGCGACCGGCAGCTCATC CTGGATGACATGCTGTGTGCCGGCAGCGAGGGCCGAGACTCCTGCTACGGTGACTCCGGCGGCGCCTCTGGTCT GCAGGCTGCGGGGGTCCTGGCGCCTGGTGGGGGTCAGCTGGGGCTACGGCTGTACCCTGCGGGACTTTC TGTGGGGGGCCACAATGCCCCCCCGGGGAAGTGGCCGTGGCAGGTCAGCCTGAGGGTCTACAGCTACCACTG CCAGCTGCCGGGGTCACCTCTCCCCAGAGTCGCTGCCGCCGCCTACCGCCTGCAGCAGGCGAGTGTGCA TGCTGAACGTCAGCCGGATCATCGTCCACCCCAACTATGTCACTGCGGGGGTGGGGTGCGGATGTGGCCCTGCT GGCCTCCTGGGCGCACATCTGTGGGGGCTCCCTCATCCACCCCCAGTGGGTGCTGACTGCTGCCCACTGCATT CCGGCGTCTACACCCACGTCCAGATCTACGTGCTCTGGATCCTGCAGCAAGTCGGGGAGTTGCCCTGA

>SGPR504_SEQID_57 ATCGGGGGCCACGAGGTGACCCCCCCACTCCAGGCCCTACATGGCATCCGTGCGCTTCGGGGGGCCAACATCACT GCGGAGGCTTCCTGCTGCGAGCCCGCTGGGTGTCTCGGCCGCCCAGTGCTTCAGCCACAGG

>SGPR469_SEQID_58

GGAGATTČTGGGĞGGCCCCTGGTCTGTGAATTAAATGGCACATGGGTCCAGGTGGGGATTGTGAGCTGGGGCA

>SGPR400 SEQID 59

CCCCAGTTTCTGTTAGCCCTGTTGTCCTGCTCATCTGCCTTCCCTCATCTGAAGTCTACCTGAAGAAGAATACAAC CAGCATCCGCCAGGGCTTGATTCACGTCTGCTCAGATACCCTCATCTCAGAGGAGTGGGTGCTGACAGTGGCGA GTGGGCGGCCCACAGTCTCATCTGGTATTGCCTCAGGCTTGGGGGCTAGTGTGGGGGCAGTGGCCCTGGCAGGT CACGGGCACTGCTGGCTATGCATTCACGCTGCTCCTTCTGCTGGGGATTTCGGGTGAGCCCCCCAGAATGGGTCT ATGGCAGĞAGAACAGGTCACCGCCAATGTCAGCAGATACCCTGGACAGAAACGATGTCCTTTCCTGAAAAAC GAAAGTGCCCCTCATTGATCTCCAGACATGCGGTGACCACTATCAAAATGAAATCTTGCTGCACGGAGTTGAGCT GAGAGTCAGGTTCTTCCCGATTTGACAGCAGGGGACGCCGCAGACCCCCCCAATTCCTCCCTTGGGTCCTGGAG TCTGCTTCCCATTATCCCCCCCACCCTGATTTCCAAGCAAACACATCTAGTGCCATCGCTGTGGTAGAACTGCCCT CTCCTGCTGGGTGACTGGATGGGGCTATACTGGAATATTCCAATATATCAAGCGTTCTTATACACTGAAGGAGCT ATTTCTCCTTTCTTATAGAGCATCACTCCTTGCTGTTGTAACACACAGATCCAATAATAGTCGTGGGCGAGCTTTT CATCATCAGTGAAGCTATGATCTGCTCCAAGCTCCCAGTGGGCCAGATGGATCAGTGTACTGTAAGAATCCACC CCTCAGGCACCTTTCACAGGCCTTGCCTTCCCCAGTGA

>SGPr397 SEQID 60

GLIHMFSIGRSYEGRCLFILKLGRRSRLKRAVWIDCGIHAREWIGPAFCQWFVKEALLTYKSDPAMRKMLNHLYFYIMP MKCLGKRRRGQAAAFLPLCWLFLKILQPGHSHLYNNRYAGDKVIRFIPKTEEEAYALKKISYQLKVDLWQPSSISYVSEG VFNVDGYHFSWTNDRFWRKTRSRNSRFRCRGVDANRNWKVKWCGKFGTNWDPDPKVSAGFTLQNMSPEDSHGR TVTDVHIPQNGSRALLAFLQEANIQYKVLIEDLQKTLEKGSSLHTQRNRRSLSGYNYEVYHSLEEIQNWMHHLNKTHS

>SGPr413_SEQID_61

WTTDRLWRKSRSPHNNGTCFGTDLNRNFNASWCSIGASRNCQDQTFCGTGPVSEPETKAVASFIESKKDDILCFLTM MKPLLETLTLGMLVPGGLGYDRSLAQHRQEIVDKSVSPWSLETYSYNIYHPMGEIYEWMREISEKYKEVVTQHFLGV HSYGQLILTPYGYTKNKSSNHPEMIQVGQKAANALKAKYGTNYRVGSSADILYASSGSSRDWARDIGIPFSYTFELRD TYETHPMYYLKISQPSGNPKKIIWMDCGIHAREWIAPAFCQWFVKEILQNHKDNSSIRKLLRNLDFYVLPVLNIDGYIYT SGTYGFVLPEAQIQPTCEETMEAVLSVLDDVYAKHWHSDSAGRVTSATMLLGLLVSCMSLL

>SGPr404_SEQID_62

RNEMTTTDDLDFKHHNYKEMRQLMKVVNEMCPNITRIYNIGKSHQGLKLYAVEISDHPGEHEVGEPEFHYIAGAHGNE LLWEAEDRQNVPRKVPNHYIAIPEWFLSENATVAAETRAVIAWMEKIPFVLGGNLQGGELVVAYPYDLVRSPWKTQEH YPHESQLPEEWENNRESLIVFMEQVHRGIKGLVRDSHGKGIPNAIISVEGINHDIRTANDGDYWRLLNPGEYVVTAKAE VLGRELLLLLVQFVCQEYLARNARIVHLVEETRIHVLPSLNPDGYEKAYEGGSELGGWSLGRWTHDGIDINNNFPDLNT MVSNDSHTWVTVKNGSGDMIFEGNSEKEIPVLNELPVPMVARYIRINPQSWFDNGSICMRMEILGCPLPDPNNYYHR TPTPDDHVFRWLAYSYASTHRLMTDARRRVCHTEDFQKEEGTVNGASWHTVAGSLNDFSYLHTNCFELSIYVGCDK **GFTASTKNCMVGYDMGATRCDFTLSKTNMARIREIMEKFGKQPVSLPARRLKLRGRKRRQRG**

GSWAFSTAAVASDRVSIHSLGHMTPVLSPQNLLSCDTHQQQGCRGGRLDGAWWFLRRRGVVSDHCYPFSGRERDE >SGPr536_1_SEQID_63 MWRCPLGLLLLPLAGHLALGAQQGRGRRELAPGLHLRGIRDAGGRYCQEQDLCCRGRADDCALPYLGAICYCDLF CNRTVSDCCPDFWDFCLGVPPFFPPIQGCMHGGRIYPVLGTYWDNCNRCTCQENRQWQCDQEPCLVDPDMIKAIN QGNYGWQAGNHSAFWGMTLDEGIRYRLGTIRPSSSVMNMHEIYTVLNPGEVLPTAFEASEKWPNLIHEPLDQGNCA AGPAPPCMMHSRAMGRGKRQATAHCPNSYVNNNDIYQVTPVYRLGSNDKEIMKELMENGPVQALMEVHEDFFLYK GGIYSHTPVSLGRPERYRRHGTHSVKITGWGEETLPDGRTLKYWTAANSWGPAWGERGHFRIVRGVNECDIESFVL GVWGRVGMEDMGHH

KYFMSPTLTMRLAGLSQITNQLHTFNDVCNNESLVSDTETSIAKELADWLISNNVVEHIFGPNLHIEIIKQCQVILNFLAAE ATHIAQGSQESCITRTGDFLGETIGNELFNCRQFIGPQHHHHHHHHHHHHDGHMVDDMLSADDVSCSSSQVSAKSEK MLMAFQNISDEQSFKAQSDHRSRHEVSHYSMWLLVSWAHCCSLVKSSLADSDHLQDWLKKLTLLIPETAVRHESCSG DIVQDEDAVNLSEGLINEAEKLLCSLVCWFTDRQIRMRFIEGCLENLGNNRSVVISLRLLPKLFGTFQQFGSSYDTHWIT LCVAKIFQIQFPLYTAYKHNTHPTIEDISTQESNILGAFCDMNDVEVPLHLLRYVCLFCGKNGLSLMKDCFEYGTPETLP GRLSTQHIDCIWAAAQLKHCSRYIHDLFPSLIKNLDPVPLRHLLNLVSALEPSVHTEQTLYLASMLIKALWNNALAAKAQ LSKQSSFASLLNTNIPIGNKKEEEELRRTAPSPWSPAASPQSSDNSDTHQSGGSDIEMDEQLINRTKHVQQRLSDTEE SMQGSSDETANSGEDGSSGPGSSSGHSDGSSNEVNSSHASQSAGSPGSEVQSEDIADIEALKEEDEDDDHGHNPP KSSCGTDLRNRKLESQAGICLGDSQGTSERNGTSSGTGKDLVFNTESLPSVDNRMRMLDACSHSEDPEHDISGEMN EYDLIGVTVHTGTADGGHYYSFIRDIVNPHAYKNNKWYLFNDAEVKPFDSAQLASECFGGEMTTKTYDSVTDKFMDFS QLCKHCTTINIDSTWQDESNQAEEPLNIDRECNEGSTERQKSIEKKSNSTRICNLTEEESSKSSDPFSLWSTDEKEKLL FLIAHAFITVVSNIRIWLHIPAVMQHIIPFRTYVIRYLCKLSDQELRQSAARNMADLMWSTVKEPLDTTLCFDKESLDLAF NMADFDGEESGCEEELVQINSHAELTSHLQQHLPNLASIYHEHLSQGPVVHKHQFNSNAVTDINLDNVCKKGNTLLW MWAEKELNIMMKLFFDNLVYYIQTVREGRQKHALYSHSAEVQVRLQFLTCVFSTLGSPDHFRLSLEQVDILWHCLVED DALARHLADCIRSREILDHQDGNVEDDGLTGLLRLATSVVKHKPPFKFSREGQEFLRDIFNLLFLLPSLKDRQQPKCKS HSSRAAAYDLLVEMVKGSVENYRLIHNWVMAQHMQSHAPYKWDYWPHEDVRAECRFVGLTNLGATCYLASTIQQLY EMSPELKNTVKSLFGGVITNNVVSLDCEHVSQTAEEFYTVRCQVADMKNIYESLDEVTIKDTLEGDNMYTCSQCGKKV VIQRLRPVHAHLYLQPGMEDGSDDMDTSVEDIGGRSCVTRFVRTLLLIMEHGVKPHSKHLTEYFAFLYEFAKMGEEES SECYDDALHWFLNQVRSKDQHAMGMETYKHLFLEKMPQLKPETISMTGLNLFQHLCNLARLATSAYDGCSNSELCG NQQAQLQEFGQSNRKGEFPGGLMGPVRMISSGHELTTDYDEKALHELGFKDMQMVFVSLGAPRRERKGEGVQLPA MCENCADLVEVLNEISDVEGGDGLQLRKEHTLKIFTYINSWTQRQCLCCFKEYKHLEIFNQVVCALINLVIAQVQVLRD MDQFWGIALRAQSGDVSRAAIQYINSYYINGKTGLEKEQEFISKCMESLMIASSSLEQESHSSLMVIERGLLMLKTHLE LYKLSLSGLDGGDSINRSFLLLAASTLLKFLPDAQALKPIRIDDYEEEPILKPGCKEYFWLLCKLVDNIHIKDASQTTLLDL RAEKRACFKKLPRIXSFNTMRYTFNMVTMMKEKVNTHFSFPLRLDMTPYTEDFLMGKSERKEGFKEVSDHSKDSESY FEKTHSAYMLFYKRMEPEEENGREYKFDVSSELLEWIWHDNMQFLQDKNIFEHTYFGFMWQLCSCIPSTLPDPKAVS AFRRFAYHLRQWQIEGTGISSHLKALSDKQSLPLRVVCQPAGLPDKMTIEMYPSDQVADLRAEVTHWYENLQKEQI MIPEARQAVFTAKYSEDMKHKTTLLELQKMFTYLMESECKAYNPRPFCKTYTMDKQPLNTGEQKDMTEFFTDLITKIE LMTAKLSTSFVLETFIHSKEKPTMLQWIELLTKQFNNSQAACEWFLDRMADDDWWPMQILIKCPNQIVRQMFQRLCIH QFLLSLQAISTMVHFYMGTKGPENPQVEVLSEEEGGEEEEEEDILSLAEEKYRPAALEKMIALVALLVEQSRSERHLTL SCLPPPQKDNIPMLLLLQEPHLTTLFDLLEMLASFKPPSGKVAVDDSESLRCEELHLHAENLSRRVWELLMLLPTCPN SQTDMAALTGGKGFPFLFQHIRDGINIRQTCNLIFSLCRYNNRLAEHIVSMLFTSIAKLTPEAANPFFKLLTMLMEFAGG

>SGPr414 SEQID 64

LQRKDVKQALIQWQERIEFAHKLLTLLNSYSPPELRNACIDVLKELVLLSPHDFLHTLVPFLQHNHCTYHHSNIPMSLGP LQEQEAKERKTKDDEGATPIKRRRVSSDEEHTVDSCISDMKTETREVLTPTSTSDNETRDSSIIDPGTEQDLPSPENSS PPGMPPFASYILQRIWEVIEYNPSQCLDWLAVQTPRNKLAHSWVLQNMENWVERFLLAHNYPRVRTSAAYLLVSLIPS TYFMDLWNLFQPKLSEPAIATNHNKQALLSFWYNVCADCPENIRLIVQNPVVTKNIAFNYILADHDDQDVVLFNRGMLP ANCANLISTLITNLISQYQNLQSDFSNRVEISKASASLNGDLRALALLLSVHTPKQLNPALIPTLQELLSKCRTCLQQRNS AYYGILRLCCEQSPAFTRQLASHQNIQWAFKNLTPHASQYPGAVEELFNLMQLFIAQRPDMREEELEDIKQFKKTTISC LVAYEGLPLHLALFPKLWTELCQTQSAMSKNCIKLLCEDPVFAEYIKCILMDERTFLNNNIVYTFMTHFLLKVQSQVFSE YFPCRENIKLIGGKSNIRPPRPELNMCLLPTMVETSKGKDDVYDRMLLDYFFSYHQFIHLLCRVAINCEKFTETLVKLSV NSFRQMFRSTRSLHIPTRDLPLSPDTTVVLHQVYNVLLGLLSRAKLYVDAAVHGTTKLVPYFSFMTYCLISKTEKLMFS VKEYRMEVPSSFSEDMSNIRSQHAEEQSNNGRYDDCKEFKDLHCSKDSTLAEEESEFPSTSISAVLSDLADLRSCDG YLRCLDGRSCWTTLISAFRILLESDEDRLLVVFNRGLILMTESFNTLHMMYHEATACHVTGDLVELLSIFLSVLKSTRPY QALPSQDPEVALSLSCGHSRGLFSHMQQHDILDTLCRTIESTIHVVTRISGKGNQAAS

>SGPr430 SEQID 65

FLSIDKVPSKDAEEMRLFLDAVHQNRLPAAMKPSQGSGSFGAILGSRTSQKETSRQLSYSDNQASAKRGSLETKDDIP FRKVLGNPGRGSIKTVAGSGIARTIPSLTSTSTPLRSGLLENRTEKRKRMISTGSELNEDYPKENDSSSNNKAMTDPSR KFNRLPRVLILHLKRYSFNVALSLNNKIGQQVIIPRYLTLSSHCTENTKPPFTLGWSAHMAMSRPLKASQMVNSCITSPS MSPLKIHĞPIRIRSMQTGITKWKEGSFEIVEKENKVSLVVHYNTGGIPRIFQLSHNIKNVVLRPSGAKQSRLMLTLQDNS LVKKDICNSETKKDLLKKVKNAISATAERFSGYMQNDAHEFLSQCLDQLKEDMEKLNKTWKTEPVSGEENSPDISATR AYTCPVITNLEFEVQHSIICKACGEIIPKREQFNDLSIDLPRRKKPLPPRSIQDSLDLFFRAEELEYSCEKCGGKCALVRH DVFDMEYTEAEAEELKRNAETGNLPHSYRLISVVSHIGSTSSSGHYISDVYDIKKQAWFTYNDLEVSKIQEAAVQSDRD KYLTSSREKQLSLKQSEENRTSGGLLPLQSSSFYGSRAGSKEHSSGGTNLDRTNVSSQTPSAKRSLGFLPQPVPLSV _AAVLEISKRDASPSLSHEDDDKPTSSPDTGFAEDDIQEMPENPDTMETEKPKTITELDPASFTEITKDCDENKENKTP PSKKFTFKSKSSLALCLDSDSEDELKRSVALSQRLCEMLGNEQQQEDLEKDSKLCPIEPDKSELENSGFDRMSEEEL KKLRCNQDYTGWNKPRVPLSSHQQQQLQGFSNLGNTCYMNAILQSLFSLQSFANDLLKQGIPWKKIPLNALIRRFAHI EGSQGEVDWLQQYDMEREREEQELQQALAQSLQEQEAWEQKEDDDLKRATELSLQEFNNSFVDALGSDEDSGNE RSGYIFFYMHKEIFDELLETEKNSQSLSTEVGKTTRQAS

FNNDGVCCCLQKRGPVNITSVCVSPRTLQISVFVLSEKYEGIVKFESDELPFGVIGSNIGDAHFQEFRAGISWKPVVDP >SGPr496_1_SEQID_66 MTLLAPWYTGPMIPMDVNEPSSVTTAPTLSSSLQHISSFLATGKKLSLHFGHPRECEVTRIDDKNRRGLEDSEPGAKL DDPIPQFPDCCSSSSRIPSVSVLVAVPLVAGHKGQAFIERMLGCFKELKQELTQEGPGGGHPRSAWPPRRHAQWP

ASSKESRMNGQLQLPTNSGNNENKYSLFAVVNHQGTLESGHYTSFIRHHKDQWFKCDDAVITKASIKDVLDSEGYLLF PGRRAELKLEPEPEPVREAEQEPKQELEDENPARSGGGGNSDEVPPTLPSDPPRPPDPSPRRSRAPRRPRPRPRPQ FFLSDRHRCEMPSPELCLVCEMSSLFRELYSGNPSPHVPYKLLHLVWIHARHLAGYRQQDAHEFLIAALDVLHRHCKG LQASTSTEVSHQQCSVPGLGEKFPTWETTKPELELLGHNPRRRRITSSFTIGLRGLINLGNTCFMNCIVQALTHTPILRD KSCICHVCGTHLNRLHSCLSCVFFGCFTEKHIHEHAETKQHNLAVDLYYGGIYCFMCKDYVYDKDIEQIAKEEQGEALK TLTDCLRRFTRPEHLGSSAKIKCGSCQSYQESTKQLTMNKLPVVACFHFKRFEHSAKQRRKITTYISFPLELDMTPFM TRLRTPPQPRPRPPRPRPRRGPGGGCLDVDFAVGPPGCSHVNSFKVGENWRQELRVIYQCFVWCGTPETRKSKA PEPCEQGEEPPPVEAEEVEEAETAEKAERKVEAEAKVEGKAEAAGKAEAAGKVDATEKVETAGKVDAAGKVETAEG DDVGKAANNPNHCNCIIDQIFTGGLQSDVTCQACHGVSTTIDPCWDISLDLPGSCTSFWPMSPGRESSVNGESHIPGI YHKQVLEHESEKVKEMNTQAY

>SGPr495_SEQID_67

MRVKDPTKALPEKAKRSKRPTVPHDEDSSDDIAVGLTCQHVSHAISVNHVKRAIAENLWSVCSECLEERRFYDGQLVL LNISNNLCFLEGKHLRSYSPQNAFQTLSQSYITTSKECSIQSCLYQFTSMELLMGNNKLLCENCTKNKQKYQEETSFAE DSEPSESESASKQTGLFRSSSGSGVQPDGPLYPLSAGKLLYTKETDSGDKEMAEAISELRLSSTVTGDQDFDRENQP **ASILKAFNNPTTKTADDETRKKVKISTVKDPFIDISLPIIEERVSKPLLWGRMNKYRSLRETDHDRYSGNVTIENIHQPRA** KKVEGVYTNARKQLLISAVPAVLILHLKRFHQAGLSLRKVNRHVDFPLMLDLAPFCSATCKNASVGDKVLYGLYGIVEH QTSAFSRIMKLCEEKCETDEIQKGGKCRNLSVRGITNLGNTCFFNAVMQNLAQTYTLTDLMNEIKESSTKLKIFPSSDS FSDIWLCLKCGFQGCGKNSESQHSLKHFKSSRTEPHCIIINLSTWIIWCYECDEKLSTHCNKKVLAQIVDFLQKHASKT QLDPLVVELSRPGPLTSALFLFLHSMKETEKGPLSPKVLFNQLCQKAPRFKDFQQQDSQELLHYLLDAVRTEETKRIQ AKKHSSSKDKSQLIHDRKCIRKLSSGETVTYQKNENLEMNGDSLMFASLMNSESRLNESPTDDSEKEASHSESNVDA SGSMREGHYTAYVKVRTPSRKLSEHNTKKKNVPGLKAADSESAGQWVHVSDTYLQVVPESRALSAQAYLLFYERVL

FIG. 2D

>SGPr407 SEQID 68

PDPSNFPTFLRCLNAFSAAVFYLPQPSWHKPEGLKPAGYPRVPDIPYGSGYTLKSTTEAAGLHQSLPMVQLPLHPTKG MEYPVPYFRSPNRTLIPERIWSNPLLVLVIAYKTVSWPRQQLLAKQANKWMPFVIPSKTLPWDPLELKICYQQNRPYPS VSLRDCLSLFTKEEELELENASGTLPVTKSEVLSTSCVPFGTTQAASTVATTQPCASARLVGTFTMTLVSPLNTLRDTE KAGKLFQPDAHGFLVKKVHAPTRGIVFIMEPRQLGGKGSLSKLQPACALGGMNSGMEPQKSAPFAAGKGLAPPLPVC RKRALASCSWCLQRVTMRRVMGVQDKAGNRNQMLLLGQRPVIGDTVSNSQTTRDKACRRPPSHSVFTQSSFWACL GIELTVMKALVLDILFKASTDIILFNHDSSSGNKWRKLPEPGGLEKKHEELRLRPLKEEYHWLVLVPLKLTGSPHRWRP SALLKESELNDADWANLMWKRYLEEQEDSKMVDLFVGQMKSYLKCQACGYHSMTFKVFFFCDLSLTIPKKGFAGGK DPDLFFYGHQSYWMKAHLNDLILREGPVTQMAQSFYWGFPAGGNLSALEMLPDGPAPRTFLQKKSCLFPLFSYILLH

NLRFKLRVYKFEEELWSRAGLGKKSDNHSSRQMPWGAAGVACQHPCKLPRIVAELTPPKLSFGFLNTVQSSVLPTSL SQFFLNDSQPEEAIPPQSLLPGSPRTNSFPKDKFVPKDKLKVILSLLTMYELDRLF

RRILMGKIFRTWFEQSPIGRKKQEEPFQEKIVVKREVKKRRQELEYQVKAELESMPPRKSLRLQGLAQSTIIEIVSVQVP NIFHGQLLSQVTCLACDNKSNTIEPFWDLSLEFPERYQCSGKDIASQPCLVTEMLAKFTETEALEGKIYVCDQCNSKRR VMHHGKGFGSGHYTAYCYNSEGGFWVHCNDSKLSMCTMDEVCKAQAYILFYTQRVTENGHSKLLPPELLLGSQHPN VFCYLCDDYVLNDNATGDLKLLRRTLSAIKSQNYHCTTRSGRFLRSMGTGDDSYFLHDGAQSLLQSEDQLYTALWHR RFSSKPVVLTEAQKQLMICHLPQVLRLHLKRFRWSGRNNREKIGVHVGFEEILNMEPYCCRETLKSLRPECFIYDLSAV >SGPr453_SEQID_69 MLAMDTCKHVGQLQLAQDHSSLNPQKWHCVDCNTTESIWACLSCSHVACGRYIEEHALKHFQESSHPVALEVNEMY MTASEKTRSCKHPPVTDTVVYQMNECQEKDTGFVCSRQSSLSSGLSGGASKGRKMELIQPKEPTSQYISLCHELHTL FQVMWSGKWALVSPFAMLHSVWRLIPAFRGYAQQDAQEFLCELLDKIQRELETTGTSLPALIPTSQRKLIKQVLNVVN AQTPASPAKDKVLSTSENEISQKVSDSSVKRRPIVTPGVTGLRNLGNTCYMNSVLQVLSHLLIFRQCFLKLDLNQWLA

>SGPr445_SEQID_70

MRVKDPTKALPEKAKRSKRPTVPHDEDSSDDIAVGLTCQHVSHAISVNHVKRAIAENLWSVCSECLEERRFYDGQLVL QTSAFSRIMKLCEEKCETDEIQKGGKCRNLSVRGITNLGNTCFFNAVMQNLAQTYTLTDLMNEIKESSTKLKIFPSSDS TSDIWLCLKCGFQGCGKNSESQHSLKHFKSSRTEPHCIIINLSTWIIWCYECDEKLSTHCNKKVLAQIVDFLQKHASKT **QLDPLVVELSRPGPLTSALFLFLHSMKETEKGPLSPKVLFNQLCQKRVHLHL**I

>SGPr401_1_SEQID_71

MTVRNIASICNMGTNASALEKDIGPEQFPINEHYFGLVNFGNTCYCNSVLQALYFCRPFRENVLAYKAQQKKKENLLTC EAQKRMRVKKLPMVLALHLKRFKYMEQLRRYTKLSYRVVFPLELRLFNTSSDAVNLDRMYDLVAVVVHCGSGPNRGH LADLFHSIATQKKKVGVIPPKKFISRLRKENDLFDNYMQQDAHEFLNYLLNTIADILQEEKKQEKQNGKLKNGNMNEPA ENNKPELTWYHEIFQGTLTNETRCLNCETVSSKDEDFLDLSVDVEQNTSITHCLRDFSNTETLCSEQKYYCETCCSKQ YITIVKSHGFWLLFDDDIVEKIDAQAIEEFYGLTSDISKNSESGYILFYQSRE

>SGPr408 SEQID 72

SDYVSQSYSYSSILNKSETGYVGLVNQAMTCYLNSLLQTLFMTPEFRNALYKWEFEESEEDPVTSIPYQLQRLFVLLQT MVPGEENÄLVPKEAPLDHTSDKSLLDANFEPGKKNFLHLTDKDGEQPQILLEDSSAGEDSVHDRFIGPLPREGSVGST SKKRAIETTDVTRSFGWDSSEAWQQHDVQELCRVMFDALEQKWKQTEQADLINELYQGKLKDYVRCLECGYEGWRI

GLEKNSLIYELFSVMAHSGSAAGGHYYACIKSFSDEQWYSFDDQHVSRITQEDIKKTHGGSSGSRGYYSSAFASSTNA VNNDRSTSSVDSDILSSSHSSDTLCNADNAQIPLANGLDSHSITSSRRTKANEGKKETWDTAEEDSGTDSEYDESGKS KTWKNPGTVFLDYHIYEEDINISSNWEVFLEVLDGVEKMKSMSQLAVLSRRWKPSEMKLDPFQEVVLESSSVDELREK DTYLDIPLVIRPYGSSQAFASVEEALHAFIQPEILDGPNQYFCERCKKKCDARKGLRFLHFPYLLTLQLKRFDFDYTTMH YMLIYRLKDPARNAKFLEVGEYPEHIKNLVQKERELEEQEKRQREIERNTCKIKLFCLHPTKQVMMENKLEVHKDKTLK FVESSETLDYQMAFADSHLWKLLDRHANTIRLFVLLPEQSPVSYSKRTAYQKAGGDSGNVDDDCERVKGPVGSLKSV EAVEMAYKMMDLEEVIPLDCCRLVKYDEFHDYLERSYEGEEDTPMGLLLGGVKSTYMFDLLLETRKPDQVFQSYKPG EVMVKVHVVDLKAESVAAPITVRAYLNQTVTEFKQLISKAIHLPAETMRIVLERCYNDLRLLSVSSKTLKAEGFFRSNKV EAILEESTEKLKSLSLQQQQDGDNGDSSKSTETSDFENIESPLNERDSSASVDNRELEQHIQTSDPENFQSEERSDSD LSEISGIPLDDIEFAKGRGTFPCDISVLDIHQDLDWNPKVSTLNVWPLYICDDGAVIFYRDKTEELMELTDEQRNELMKK RIKLNDRMTFPEELDMSTFIDVEDEKSPQTESCTDSGAENEGSCHSDQMSNDFSNDDGVDEGICLETNSGTEKISKS RGEMQYMYFKAEPYAADEGSGEGHKWLMVHVDKRITLAAFKQHLEPFVGVLSSHFKVFRVYASNQEFESVRLNETL SSFSDDNKITIRLGRALKKGEYRVKVYQLLVNEQEPCKFLLDAVFAKGMTVRQSKEELIPQLREQCGLELSIDRFRLRK **ESSRLQKTGHRVTYSPRKEKALKIYLDGAPNKDLTQD**

>SGPr480_SEQID_73

NLIVGLVLLTRGKDEEKAKYIFSLFSSESGNYVIREEMERMLHVVDGKVPDTLRKCFSEGEKVNYEKFRNWLFLNKDAF GHNWFIISMQWWQQWKEYVKYDANPVVIEPSSVLNGGKYSFGTAAHPMEQVEDRIGSSLSYVNTTEEKFSDNISTAS KRTPQLIHGRDYEMVPEPVWRALYHWYGANLALPRPVIKNSKTDIPELELFPRYLLFLRQQPATRTQQSNIWVNMGNV PSPNAPLKRVLAYTGCFSRMQTIKEIHEYLSQRLRIKEEDMRLWLYNSENYLTLLDDEDHKLEYLKIQDEQHLVIEVRNK DMSWPEEMSFIANSSKIDRHKVPTEKGATGLSNLGNTCFMNSSIQCVSNTQPLTQYFISGRHLYELNRTNPIGMKGHM MGAKESRIGFLSYEEALRRVTDVELKRLKDAFKRTCGLSYYMGQHCFIREVLGDGVPPKVAEVIYCSFGGTSKGLHFN EASETAGSGFLYSATPGADVCFARQHNTSDNNNQCLLGANGNILLHLNPQKPGAIDNQPLVTQEPVKATSLTLEGGRI DWÉVAAEAWDNHLRRNRSIVVDLFHGQLRSQVKCKTCGHISVRFDPFNFLSLPLPMDSYMHLEITVIKLDGTTPVRYG PWYRFCRGCKIDCGEDRAFIGNAYIAVDWDPTALHLRYQTSQERVVDEHESVEQSRRAQAEPINLDSCLRAFTSEEEL SPSTNEMFTLTTNGDLPRPIFIPNGMPNTVVPCGTEKNFTNGMVNGHMPSLPDSPFTGYIIAVHRKMMRTELYFLSSQ KNRPSLFGMPLIVPCTVHTRKKDLYDAVWIQVSRLASPLPPQEASNHAQDCDDSMGYQYPFTLRVVQKDGNSCAWC AKCYGDLVQELWSGTQKNVAPLKLRWTIAKYAPRFNGFQQQDSQELLAFLLDGLHEDLNRVHEKPYVELKDSDGRP TFSRWLLSGGVYVTLTDDSDTPTFYQTLAGVTHLEESDIIDLEKRYWLLKAQSRTGRFDLETFGPLVSPPIRPSLSEGL FNAFDENRDNHIDFKEISCGLSACCRGPLAERQKFCFKVFDVDRDGVLSRVELRDMVVALLEVWKDNRTDDIPELHM DLSDIVEGILNAHDTTKMGHLTLEDYQIWSVKNVLANEFLNLLFQVCHIVLGLRPATPEEEGQIIRGWLERESRYGLQA LRLNMDEKYTGLKKQLSDLCGLNSEQILLAEVHGSNIKNFPQDNQKVRLSVSGFLCAFEIPVPVSPISASSPTQTDFSS

KGRLRLPQIGSKNKLSSSKENLDASKENGAGQICELADALSRGHVLGGSQPELVTPQDHEVALANGFLYEHEACGNG YSNGQLGNHSEEDSTDDQREDTRIKPIYNLYAISCHSGILGGGHYVTYAKNPNCKWYCYNDSSCKELHPDEIDTDSAY QGDELSEPRILAREVKKVDAQSSAGEEDVLLSKSPSSLSANIISSPKGSPSSSRKSGTSCPSSKNSSPNSSPRTLGRS GENEMYYCSKCKTHCLATKKLDLWRLPPILIIHLKRFQFVNGRWIKSQKIVKFPRESFDPSAFLVPRDPALCQHKPLTP ILFYEQQGIDYAQFLPKTDGKKMADTSSMDEDFESDYKKYCVLQ

>SGPr431_SEQID_74

SETSLQEVASKAAVLTETPRTSDGEKTLIEKMFGGKLRTHIRCLNCRSTSQKVEAFTDLSLAFCPSSSLENMSVQDPAS MDKILEGLVSSSHPLPLKRVIVRKVVESAEHWLDEAQCEAMFDLTTRLILEGQDPFQRQVGHQVLEAYARYHRPEFES PKGKLSITFCQQLVRTIGHFQCVSTQERELREYVSQVTKVSNLLQNIWKAEPATLLPSLQEVFASISSTDASFEPSVALA NGCNSLMKKLQHLFAFLAHTQREAYAPRIFFEASRPPWFTPRSQQDCSEYLRFLLDRLHEEEKILKVQASHKPSEILEC TPSVTDLLNYFLAPEILTGDNQYYCENCASLQNAEKTMQITEEPEYLILTLLRFSYDQKYHVRRKILDNVSLPLVLELPVK FFNKTFVLGLLHQGYHSLDRKDVAILDYIHNGLKLIMSCPSVLDLFSLLQVEVLRMVCERPEPQLCARLSDLLTDFVQCI SLVQHIPLQMITVLIRSLTTDPNVKDASMTQALCRMIDWLSWPLAQHVDTWVIALLKGLAAVQKFTILIDVTLLKIELVFN RLWFPLVRPGALAVLSHMLLSFQHSPEAFHLIVPHVVNLVHSFKNDGLPSSTAFLVQLTELIHCMMYHYSGFPDLYEPI RITSFSSLSESWSVDVDFTDLSENLAKKLKPSGTDEASCTKLVPYLLSSVVVHSGISSESGHYYSYARNITSTDSSYQM LEAIKDFPKPSEEKIKLILNQSAWTSQSNSLASCLSRLSGKSETGKTGLINLGNTCYMNSVIQALFMATDFRRQVLSLNL YHQSEALALASSQSHLLGRDSPSAVFEQDLENKEMSKEWFLFNDSRVTFTSFQSVQKITSRFPKDTAYVLLYKKQHS1 NGLSGNNPTSGLWINGDPPLQKELMDAITKDNKLYLQEQELNARARALQAASASCSFRPNGFDDNDPPGSCGPTGG SPSIQDGGLMQASVPGPSEEPVVYNPTTAAFICDSLVNEKTIGSPPNEFYCSENTSVPNESNKILVNKDVPQKPGGET GGGGFNTVGRLVF

>SGPr429_SEQID_75

PSKSQTRKPKCGKGTHCSRNAYMLVYRLQTQEKPNTTVQVPAFLQELVDRDNSKFEEWCIEMAEMRKQSVDKGKAK HEEVKELYQRLPAGAEPYEFVSLEWLQKWLDESTPTKPIDNHACLCSHDKLHPDKISIMKRISEYAADIFYSRYGGGPR MAPRLQLEKAAWRWAETVRPEEVSQEHIETAYRIWLEPCIRGVCRRNCKGNPNCLVGIGEHIWLGEIDENSFHNIDDP NSNRRYIDPSGFVKALGLDTGQQQDAQEFSKLFMSLLEDTLSKQKNPDVRNIVQQQFCGEYAYVTVCNQCGRESKLL SKFYELELNIQGHKQLTDCISEFLKEEKLEGDNRYFCENCQSKQNATRKIRLLSLPCTLNLQLMRFVFDRQTGHKKKLN TYIGFSEILDMEPYVEHKGGSYVYELSAVLIHRGVSAYSGHYIAHVKDPQSGEWYKFNDEDIEKMEGKKLQLGIEEDLE NCERRKKNSFVGLTNLGATCYVNTFLQVWFLNLELRQALYLCPSTCSDYMLGDGIQEEKDYEPQTICEHLQYLFALLQ LTVKALCKECVVERCRILRLKNQLNEDYKTVNNLLKAAVKGDGFWVGKSSLRSWRQLALEQLDEQDGDAEQSNGKM NGSTLNKDESKEERKEEEELNFNEDILCPHGELCISENERRLVSKEAWSKLQQYFPKAPEFPSYKECCSQCKILEREG

GLMFTFASMTKEDSKLIALIWPSEWQMIQKLFVVDHVIKITRIEVGDVNPSETQYISEPKLCPECREGLLCQQQRDLREY RHRKVRGEKALLVSANQTLKELKIQIMHAFSVAPFDQNLSIDGKILSDDCATLGTLGVIPESVILLKADEPIADYAAMDDV EENEALHKMIANEQKTSLPNLFQDKNRPCLSNWPEDTDVLYIVSQFFVEEWRKFVRKPTRCSPVSSVGNSALLCPHG TQATIYVHKVVDNKKVMKDSAPELNVSSSETEEDKEEAKPDGEKDPDFNQSNGGTKRQKISHQNYIAYQKQVIRRSM MQVCMPEEGFKGTGLLGH

>SGPr503 SEQID 76

MNSLLQGLSACPAFIRWLEEFTSQYSRDQKEPPSHQYLSLTLLHLLKALSCQEVTDDEVLDASCLLDVLRMYRWQISS CKHCEHQSPVRFDTFDSLSLSIPAATWGHPLTLDHCLHHFISSESVRDVVCDNCTKIEAKGTLNGEKVEHQRTTFVKQ PVLNQPGAPKTQIFMNGACSPSLLPTLSAPMPFPLPVVPDYSSSTYLFRLMAVVVHHGDMHSGHFVTYRRSPPSARN MLSSRAEAAMTAADRAIQRFLRTGAAVRYKVMKNWGVIGGIAAALAAGIYVIWGPITERKKRRKGLVPGLVNLGNTCF LKLGKLPQCLCIHLQRLSWSSHGTPLKRHEHVQFNEFLMMDIYKYHLLGHKPSQHNPKLNKNPGPTLELQDGPGAPT FEEQDAHELFHVITSSLEDERDRQPRVTHLFDVHSLEQQSEITPKQITCRTRGSPHPTSNHWKSQHPFHGRLTSNMV PLSTSNQWLWVSDDTVRKASLQEVLSSSAYLLFYERVLSRMQHQSQECKSEE

>SGPr427 SEQID 77

LREDEVNTRGAYILFYQKRNSIPPWSASSSMRGSTSSSLSDHWLLRLGSHAGSTRGSLLSWSSAPCPSLPQVPDSPIF MDLGPGDÄAGGGPLAPRPRRRRSLRRLFSRFLLALGSRSRPGDSPPRPQPGHCDGDGEGGFACAPGPVPAAPGSP RALWTREYTPQLSAEFKNAVSKYGSQFQGNSQHDALEFLLWLLDRVHEDLEGSSRGPVSEKLPPEATKTSENCLSPS **ALRKMVAEEGGVPADEVILVELYPSGFQRSFFDEEDLNTIAEGDNVYAFQVPPSPSQGTLSAHPLGLSASPRLAAREG** SVACSYLSPKDSRPLCHWAVDRVLHLRRPGGPPHVKLAVEWDSSVKERLFGSLQEERAQDADSVWQQQQAHQQH SCTLDECFQFYTKEEQLAQDDAWKCPHCQVLQQGMVKLSLWTLPDILIIHLKRFCQVGERRNKLSTLVKFPLSGLNMA SQSIVSLLTGTAGEDEKSASPRSNVALPANSEDGGRAIERGPAGVPCPSAQPNHCLAPGNSDGPNTARKLKENAGQD GEERPPGPQPQLQLPAGDGARPPGAQGLKNHGNTCFMNAVVQCLSNTDLLAEFLALGRYRAAPGRAEVTEQLAALV PHVAQRSTSPEAGLGPWPSWKQPDCLPTSYPLDFLYDLYAVCNHHGNLQGGHYTAYCRNSLDGQWYSYDDSTVEP AQLPLGQSFVQSHFQAQYRSSLTCPHCLKQSNTFDPFLCVSLPIPLRQTRFLSVTLVFPSKSQRFLRVGLAVPILSTVA QRFSLSLHSESKVLILFCNLVGSGQQASRFGPPFLIREDRAVSWAQLQQSILSKVRHLMKSEAPVQNLGSLFSIRVVGL IKLPRKFDLPLTVMPSVEHEKPARPEGQKAMNWKESFQMGSKSSPPSPYMGFSGNSKDSRRGTSELDRPLQGTLTL TNSLCNQEKGGLEPRRLVRGVKGRSISMKAPTTSRAKQGPFKTMPLRWSFGSKEKPPGASVELVEYLESRRRPRST LRSVFRKKENRRNERAEVSPQVPPVSLVSGGLSPAMDGQAPGSPPALRIPEGLARGLGSRLERDVWSAPSSLRLPR KASRAPRGSALGMSQRTVPGEQASYGTFQRVKYHTLSLGRKKTLPESSF

FIG. 21

WAHEDGWPFDGPGGILGHAFLPNSGNPGVVHFDKNEHWSASDTGYNLFLVATHEIGHSLGLQHSGNQSSIMYPTYW HQPHCGVPDGSDTSISPGRCKWNKHTLTYRIINYPHDMKPSAVKDSIYNAVSIWSNVTPLIFQQVQNGDADIKVSFWQ RHVQQIDAAVYLREPQKTLFFVGDEYYSYDERKRKMEKDYPKNTEEEFSGVNGQIDAAVELNGYIYFFSGPKTYKYDT GLAHSTDPSALMYPTYKYKNPYGFHLPKDDVKGIQALYGPRKVFLGKPTLPHAPHHKPSIPDLCDSSSSFDAVTMLGK FGLQVTGKLDQTTMNVIKKPRCGVPDVANYRLFPGEPKWKKNTLTYRISKYTPSMSSVEVDKAVEMALQAWSSAVPL SFVRINSGEADIMISFENGDHGDSYPFDGPRGTLAHAFAPGEGLGGDTHFDNPEKWTMGTNGFNLFTVAAHEFGHAL MKVLPASGLAVFLIMALKFSTAAPSLVAASPRTWRNNYRLAQAYLDKYYTNKEGHQIGEMVARGSNSMIRKIKELQAF ELLLFKDRIFWRRQVHLRTGIRPSTITSSFPQLMSNVDAAYEVAERGTAYFFKGPHYWITRGFQMQGPPRTIYDFGFP MQLVILRVTIFLPWCFAVPVPPAADHKGWDFVEGYFHQFFLTKKESPLLTQETQTQLLQQFHRNGTDLLDMQMHALL YHDPRTFQLSADDIQRIQHLYGEKCSSDIP EKEDVVSVVKSSSWIGC

>SGPr092 SEQID 78

>SGPr104_1_SEQID_80 MNVALQELGAGSNMVEYKRATLRDEDAPETPVEGGASPDAMEVGFQKGTRQLLGSRTQLELVLAGASLLLAALLLGC LKHLLENTTFNSSSEAEQKTQRFYLSCLQVERIEELGAQPLRDLIEKIGGWNITGPWDQDNFMEVLKAVAGTYRATPFF EEALGQLVWMDEKTRQAAKEKADAIYDMIGFPDFILEPKELDDVYDGYEISEDSFFQNMLNLYNFSAKVMADQLRKPP WQNESLAAFRNHTACMEEQYNQYQVNGERLNGRQTLGENIADNGGLKAAYNAYKAWLRKHGEEQQLPAVGLTNHQ LVALGVQYHRDPSHSTCLTEACIRVAGKILESLDRGVSPCEDFYQFSCGGWIRRNPLPDGRSRWNTFNSLWDQNQAI TVYISADSKSSNSNVIQVDQSGLFLPSRDYYLNRTANEKVLTAYLDYMEELGMLLGGRPTSTREQMQQVLELEIQLANI NLVQKTTSSLDRRFESAQEKLLETLYGTKKSCVPRWQTCISNTDDALGFALGSLFVKATFDRQSKEIAEGMISEIRTAF TVPQDQRRDEEKIYHKMSISELQALAPSMDWLEFLSFLLSPLELSDSEPVVVYGMDYLQQVSELINRTEPSILNNYLIW SRDQWSMTPQTVNAYYLPTKNEIVFPAGILQAPFYARNHPKALNFGGIGVVMGHELTHAFDDQGREYDKEGNLRPW LFFVGFAQVWCSVRTPESSHEGLVTDPHSPARFRVLGTLSNSRDFLRHFGCPVGSPMNPGQLCEVW

>SGPr303 SEQID 81

EDEKVTLSFPSTLQTGTGTLKIDFVGELNDKMKGFYRSKYTTPSGEVRYAAVTQFEATDARRAFPCWDERAIKATFDIS MPEKRPFERLPADVSPINCSLCLKPDLLDFTFEGKLEAAAQVRQATNQIVMNCADIDIITASYAPEGDEEIHATGFNYQN ALEVAAKTLPFYNDYFNVPYPLPKIDLIAIADFAAGAMENWDLVTYRETALLIDPKNSCSSSRQWVALVVGHELAHQWF LVVPKDRVALSNMNVIDRKPYPDDENLVEVKFARTPVTSTYLVAFVVGEYDFVETRSKDGVCVCVYTPVGKAEQGKF

GNLVTMEWWTHLWLNEGFASWIEYLCVDHCFPEYDIWTQFVSADYTRAQELDALDNSHPIEVSVGHPSEVDEIFDAIS YSKGASVIRMLHDYIGDKVKKKTLSI

>SGPr402 1 SEQID 82

VVQQSLTPHWGHHLHLKKNPKVQWFQQQTLQRRVKRSVVVPTDPWFSKQWYMNSEAQPDLSILQAWSQGLSGQGI VVSVLDDGIEKDHPDLWANYDPLASYDFNDYDPDPQPRYTPSKENRHGTRCAGEVAAMANNGFCGVGVAFNARIGG VRMLDGTITDVIEAQSLSLQPQHIHIYSASWGPEDDGRTVDGPGILTREAFRRGVTKGRGGLGTLFIWASGNGGLHYD LGQLCLAYCPPRFFNHTRLVTAGPGHTAAPALRVCSSCHASCYTCRGGSPRDCTSCPPSSTLDQQQGSCMGPTTPD MRPAPIALWLRLVLALALVRPRAVGWAPVRAPIYVSSWAVQVSQGNREVERLARKFGFVNLGPIFPDGQYFHLRHRG ALALEANPFLTWRDMQHLVVRASKPAHLQAEDWRTNGVGRQVSHHYGYGLLDAGLLVDTARTWLPTQPQRKCAVR VQSRPTPILPLIYIRENVSACAGLHNSIRSLEHVQAQLTLSYSRRGDLEISLTSPMGTRSTLVAIRPLDVSTEGYNNWVF MSTHFWDENPQGVWTLGLENKGYYFNTGTLYRYTLLLYGTAEDMTARPTGPQVTSSACVQRDTEGLCQACDGPAYI NCNCDGYTNSIHTLSVGSTTQQGRVPWYSEACASTLTTTYSSGVATDPQIVTTDLHHGCTDQHTGTSASAPLAAGMI SRPRLRAAACPHHRCPASAMVLSLLAVTLGGPVLCGMSMDLPLYAWLSRARATPTKPQVWLPAGT

>SGPr434 SEQID 83

CATNYGENLCYGDSGGPLACEVEGRWILAGVLSWEKACVKAQNPGVYTRITKYTKWIKKQMSNGAFSGPCASACLLF <u> GPGRQGGCAGRRSTALPLRAPLRARRPGPRSERMGAATCRGSRIPSGPPVQGERSAPRFGVTSLSLWPADFKDNW</u> RIAGSRQEVALAGEPADQQQTHLRRLPYRQTLGYKEDTTNPVCGEPWWSEDLEMTRHWPWEVSLRMENEHVCGG QPICLPEPNFNLKVGTQCWVTGWSQVKQRFSGSTANSMLTPELQEAEVFIMDNKRCDRHYKKSFFPPVVPLVLGDMI ALIDPSWVVTAAHCSQGTKEYSVVLGTSKLQPMNFSRALWVPVRDIIMHPKYWGRAFIMGDVALVHLQTPVTFSEYV

>SGPr446_1_SEQID_84

VVSVQRAFVHPKFSTVTTIRNDLALLQLQHPVNFTSNIQPICIPQENFQVEGRTRCWVTGWGKTPERGEKLASEILQDV ILTPVCGRTPLRIVGGVDAEEGRWPWQVSVRTKGRHICGGTLVTATWVLTAGHCISSRFHYSVKMGDRSVYNENTSV DQYIMCYEECNKIIQKALSSTKDVIIKGMVCGYKEQGKDSCQGDSGGRLACEYNDTWVQVGIVSWGIGCGR

>SGPr447 SEQID 85

QKHYYPSEWTVQLGELTSRPTPWNLRAYSSRYKVQDIIVNPDALGVLRNDIALLRLASSVTYNAYIQPICIESSTFNFVH MGARGALTLALLLÄRAGLGKPEACGHREIHALVAGGVESARGRWPWQASLRLRRRHRCGGSLLSRRWVLSAAHCF

RPDCWVTGWGLISPSGTPLPPPYNLREAQVTILNNTRCNYLFEQPSSRSMIWDSMFCAGAEDGSVDTCKGDSGGPL VCDKDGLWYQVGIVSWGMDCGQPNRPGVYTNISVYFHWIRRVMSHSTPRPNPSQLLLLLALLWAP

LPGMVCTSAVGELPSCEGLSGAPLVHEVRGTWFLAGLHSFGDACQGPARPAVFTALPAYEDWVSSLDWQVYFAEEP YDMALLLLAQPVTLGASLRPLCLPYPDHHLPDGERGWVLGRARPGAGISSLQTVPVTLLGPRACSRLHAAPGGDGSPI GASCWATGWDQDTSDAPGTLRNLRLRLISRPTCNCIYNQLHQRHLSNPARPGMLCGGPQPGVQGPCQGDSGGPVL NQSNSIFGCIFYTLQLLLGLQAAQRACGQRGPGPPKPQEGNTVPGEWPWQASVRRQGAHICSGSLVADTWVLTAAH **CFEKAAATELNSWSVVLGSLQREGLSPGAEEVGVAALQLPRAYNHYSQGSDLALLQLAHPTTHTPLCLPQPAHRFPF** CLEPDGHWVQAGIISFASSCAQEDAPVLLTNTAAHSSWLQARVQGAAFLAQSPETPEMSDEDSCVACGSLRTAGPQ AGAPSPWPWEARLMHQGQLACGGALVSEEAVLTAAHCFIGRQAPEEWSVGLGTRPEEWGLKQLILHGAYTHPEGG >SGPr432_1_SEQID_86 MGSTWGSPGWVRLALCLTGLVLSLYALHVKAARARDRDYRALCDVGTAISCSRVFSSRWGRGFGLVEHVLGQDSIL **EPEAEPGSCLANISQPTSC**

DQSWVLTAAHCGNKPLWARVGDDHLLLLQGEQLRRTTRSVVHPKYHQGSGPILPRRTDEHDLMLLKLARPVVPGPR >SGPr529_SEQID_87 MRAPHLHLSAASGARALAKLLPLLMAQLWAAEAALLPQNDTRLDPEAYGAPCARGSQPWQVSLFNGLSFHCAGVLV VRALQLPYRCAQPGDQCQVAGWGTTAARRVKYNKGLTCSSITILSPKECEVFYPGVVTNNMICAGLDRGQDPCQSD SGGPLVCDETLQGILSWGVYPCGSAQHPAVYTQICKYMSWINKVIRSN

>SGPr428_1_SEQID_88

QYDDLNMHLEKLRVVQISRKECAKRVNQLSRNMICAWNEPGTNGQGPGEVGGPLVCQKKNKSTWYQLGIISWGVGC NVQCGHRPÄFPNSSWLPFHERLQVQNGECPWQVSIQMSRKHLCGGSILHWWWVLTAAHCFRRTLLDMAVVNVTVV MGTRTFSNIHSERKQVQKVIIHKYYKPPQLDSDLSLLLLATPVQFSNFKMPVCLQEEERTWDWCWMAQWVTTNGYD GQKNMPGVYTELSNYLLWIERKTVLAGKPYKYEPDSVYALLLSPWAILLLYFVMLLLS

>SGPr425 SEQID 89

EDYLSYETVFENGTRTLTRVKVQDLVLEPTQNITTKGVSVRRKRQVYGTDSRFSILDKRFLTNFPFSTAVKLSTGCSGIL MENMLLW\(\bar{U}\) IFFTP\(\bar{G}\) WTLIDGSEMEWDFMWHLRKVPRIVSERTFHLTSPAFEADAKMMVNTVCGIECQKELPTPSLSEL |SPQHVLTAAHCVHDGKDYVKGSKKLRVGLLKMRNKSGGKKRRGSKRSRREASGGDQREGTREHLPERAKGGRRR KKSGRGQRIAEGRPSFQWTRVKNTHIPKGWARGGMGDATLDYDYALLELKRAHKKKYMELGISPTIKKMPGGMIHFS

FIG. 2L

GFDNDRADQLVYRFCSVSDESNDLLYQYCDAESGSTGSGVYLRLKDPDKKNWKRKIIAVYSGHQWVDVHGVQKDYN **VAVRITPLKYAQICLWIHGNDANCAYG**

>SGPr548_SEQID_90

DTLHCANISIISDTSCDKSYPGRLTNTMVCAGAEGRGAESCEGDSGGPLVCGGILQGIVSWGDVPCDNTTKPGVYTKV TTSRVIPHPRYEARSHRNDIMLLRLVQPARLNPQVRPAVLPTRCPHPGEACVVSGWGLVSHNEPGTAGSPRSQVSLP MGDPEGSAEWGWGKGIPVVRRNLLTVDGISLCLEGSWWRQKGPASPGFSHSLPRLQPNPGPSSTMWLLLTLSFLLA STAAQDGDKLLEGDECAPHSQPWQVALYERGRFNCGASLISPHWVLSAAHCQSRFMRVRLGEHNLRKRDGPEQLR CHYLEWIRETMKRN

>SGPr396 SEQID 91

MGPAGCAFTLLLLLLGISVCGQPVYSSRVVGGQDAAAGRWPWQVSLHFDHNFICGGSLVSERLILTAAHCIQPTWTTFS VSWGLECGKSLPGVYTNVIYYQKWINATISRANNLDFSDFLFPIVLLSLALLRPSCAFGPNTIHRVGTVAEAVACIQGWE YTVWLGSITVGDSRKRVKYYVSKIVIHPKYQDTTADVALLKLSSQVTFTSAILPICLPSVTKQLAIPPFCWVTGWGKVKE SSDRDYHSALQEAEVPIIDRQACEQLYNPIGIFLPALEPVIKEDKICAGDTQNMKDSCKGDSGGPLSCHIDGVWIQTGV ENAWRFSPRGR

>SGPr426_SEQID_92

KPSFRLTRCGIRMTSSNMPLPASSSTQRIVQGRETAMEGEWPWQASLQLIGSGHQCGASLISNTWLLTAAHCFWKNK VDDGP!QNTLRQARVETISTDVCNRKDVYDGLITPGMLCAGFMEGKIDACKGDSGGPLVYDNHDIWYIVGIVSWGQSC REFIERSHQIERMMSRIFRHSSVGGRFIKSHVIKLSPDEQGVDILIVLIFRYPSTDSAEQIKKKIEKALYQSLKTKQLSLTIN MMYAPVEFSEAEFSRAEYQRKQQFWDSVRLALFTLAIVAIIGIAIGIVTHFVVEDDKSFYYLASFKVTNIKYKENYGIRSS DPTQWIATFGATITPPAVKRNVRKIILHENYHRETNENDIALVQLSTGVEFSNIVQRVCLPDSSIKLPPKTSVFVTGFGSI **ALPKKPGVYTRVTKYRDWIASKTGM**

>SGPr552 SEQID 93

RIAEGLDÄEEGEWPWQASLPQNNVYRRGATWLSNSWLITAAHCFIRVHDPKEWNVILSNPQTQSNIKNVIIQENYHYP AHDNDIAVVHLSSPVLYTSNIQKACLPDVNYIFLYNSEAVVTAWGSFKPLRTTSNVLHKGLVKIIDNRTCNNGEADGRVI TSGMLCAGFLEPRVDACQGDSGGPLVGTDSKGILAKGSLLVLKAGVNERALPNKPSVYTQVTYY

>SGPr405 SEQID 94

MVSKGGVAAEPEPHYCEDSERGPNTLTGPGSLPRGGGIEVGMEFPGCSGEGCVKPHEEAAREGAGRGKRAVPGPK GKMAEFNWSMAFKGPAAGHEERLNSVSSRAKKGIGWDVAAASLRGVDHFSDLPPPLQVREELEACAFRVQVGQLRL WPLSLWEATVKVRSNVLCNQTCRRRFPSNHTERFERLIKDDMLCAGDGNHGSWPGDNGGPLLCRRNCTWVQVEVV MRTRACERMYHKGPTAHGQVTIIKAAMPCAGRKGQGSCQAALRTEDLTPTTPNTEVSPRADPRLSQPEDIWPEWAW RRQQGSAEGPAAGWTLEQETRGDVLEDKNERADEEILRLAPGKGRLPIDSKHLKPVISSFPVRSQELGEGAGAGTLR PVVVGTTMLLLLFLAVSSLGSCSTGSPAPVPENDLVGIVGGHNTPGEVVVAVGADRRSLHFPEGHRPVHLPDSHQG YEDDQRTKVVEIVRHPQYNESLSAQGGADIALLKLEAPVPLSELIHPVSLPSASLDVPSGKTCWVTGWGVIGRGELLP SNADCERQTYDAFPGAGDRKFIQDDMICAGRTGRRTWKGDSGGPLVCKKKGTWLQAGVVSWGFYSDRPSIGVYTR PETSWQGANHADAQRPAGRVPTMQRPRDMGQGQEWVCRPFTHVTCYPTAIPRPFTHVTCYLMAVPSTLTHVTCYP FAVPRPFTHVTCYLMAVPSTLTHITCYMMAVPRPFTHITCYPMAVPSTLTHVTCHPTAIPRPFTHITCYTMAIPRPSTTP SWGKLCGLRGYPGMYTRVTSYVSWIRQPCPSAQTPAVVRRFVLPPNPDVEALTPSVMGSGAPLPPAPDLQEAEVPI CVSVRGPGAAECQPDRRPPNYSVFFLGADIALLKLATSSLEFTDSDNCWNTGWGMVGLLDMLPPPYRPQQVKVLTL PATRRPSPAPSPTSPATRWPSPGPSPMSPATR

>SGPr485_1_SEQID_95

LLLLASPIKLDDLKVPICLPTQPGPATWRECWVAGWGQTNAADKNSVKTDLMKAPMVIMDWEECSKMFPKLTKNMLC AGYKNESYDACKGDSGGPLVCTPEPGEKWYQVGIISWGKSCGEKNTPGIYTSLVNYNLWIEKVTQLEGRPFNAEKRR FPWQVSIQVRSEPFCGGSILNKWWILTAAHCLYSEELFPEELSVVLGTNDLTSPSMEIKEVASIILHKDFKRANMDNDIA MLLFSVLLĪLĪSLVTRĪQLGPRTPLPEAGVAILGRARGAHRPQPPHPPSPVSECGDRSIFEGRTRYSRITGGMEAEVGE TSVKQKPMGSPVSGVPEPGSPRSWLLLCPLSHVLFRAILY

GVRTSDVVVAGEFDQGSDEENIQVLKIAKVFKNPKFSILTVNNDITLLKLATPARFSQTVSAVCLPSADDDFPAGTLCAT TGWGKTKYNANKTPDKLQQAALPLLSNAECKKSWGRRITDVMICAGASGVSSCMGDSGGPLVCQKDGAWTLVGIVS >SGPr534_SEQID_96 MASLWLLSCFSLVGAAFGCGVPAIHPVLSGLSRIVNGEDAVPGSWPWQVSLQDKTGFHFCGGSLISEDWVVTAAHC WGSRTCSTTTPAVYARVTKLIPWVQKILAAN

>SGPr390 SEQID 97

NPPEGRNSEHIRTSARTNSGHTIFKKCNTQPFLSTQGFHVDHTAELRGIRWTSSLRRETSDYHRTLTPTLEALLHFLLR MEPTVADVHLVPRTTKEVPALDAACCRAASIGVVATSLVVLTLGVLLGGMNNSRHAALRAATLPGKVYSVTPEASKTT

DPTKWVAYVGATYLSGSEASTVRAQVVQIVKHPLYNADTADFDVAVLELTSPLPFGRHIQPVCLPAATHIFPPSKKCLIS GWGYLKEDFRKHLPRPAMVKPEVLQKATVELLDQALCASLYGHSLTDRMVCAGYLDGKVDSCQGDSGGPLVCEEPS GWGNTQEGNATKPELLQKASVGIIDQKTCSVLYNFSLTDRMICAGFLEGKVDSCQGDSGGPLACEEAPGVFYLAGIVS WGIGCAQVKKPGVYTRITRLKGWILEIMSSQPLPMSPPSTTRMLATTSPRTTAGLTVPGATPSRPTPGAASRVTGQPA DDQEDCSDGSDEAHCECGLQPAWRMAGRIVGGMEASPGEFPWQASLRENKEHFCGAAIINARWLVSAAHCFNEFQ PLQTLSLGLEEELLQRGIRARLREHGISLAAYGTIVSAELTGRHKGPLAERDFKSGRCPGNSFSCGNSQCVTKVNPEC GRFFLAGIVSWGIGCAEARRPGVYARVTRLRDWILEATTKASMPLAPTMAPAAAPSTAWPTSPESPVVSTPTKSMQ NSTLSAVSTTARGQTPFPDAPEATTHTQLPDCGLAPAALTRIVGGSAAGRGEWPWQVSLWLRRREHRCGAVLVAER WLLSAAHCFDVYGDPKQWAAFLGTPFLSGAEGQLERVARIYKHPFYNLYTLDYDVALLELAGPVRRSRLVRPICLPEP ALSTVPLDWVTVPKLQECGARPAMEKPTRVVGGFGAASGEVPWQVSLKEGSRHFCGATVVGDRWLLSAAHCFNHT KVEQVRAHLGTASLLGLGGSPVKIGLRRVVLHPLYNPGILDFDLAVLELASPLAFNKYIQPVCLPLAIQKFPVGRKCMIS APRPPDGTRCVITGWGSVREGGSMARQLQKAAVRLLSEQTCRRFYPVQISSRMLCAGFPQGGVDSCSGDAGGPLA CREPSGRWVLTGVTSWGYGCGRPHFPGVYTRVAAVRGWIGQHIQE

>SGPr521_SEQID_98

MARSLLLPLQILLLSLALETAGEEAQGDKIIDGAPCARGSHPWQVALLSGNQLHCGGVLVNERWVLTAAHCKMNEYTV HLGSDTLGDRRAQRIKASKSFRHPGYSTQTHVNDLMLVKLNSQARLSSMVKKVRLPSRCEPPGTTCTVSGWGTTTS PDVTFPSDLMCVDVKLISPQDCTKVYKDLLENSMLCAGIPDSKKNACNGDSGGPLVCRGTLQGLVSWGTFPCGQPN **DPGVYTQVCKFTKWINDTMKKHR**

FIG. 2N

>SGPr530_1_SEQID_99

HQTQHTQKMSVHRIITHPDFEKLHPFGSDIAMLQLHLPMNFTSYIVPVCLPSRDMQLPSNVSCWITGWGMLTEDHKRV VSTVCGKPKVVGKIYGGRDAAAGQWPWQASLLYWGSHLCGAVLIDSCWLVSTTHCFLNKSQAPKNYQVLLGNIQLY QLSPPFYLQEGKVGLIENTLCNTLYGQRTAKARPKLCTRRCCVGGYFSTGKSICKGDSGGPLVCYLPSAWVLVGLAS WGLDCRHPAYPSIFTRVTYFINWIDEIMRLTPLSDPALAPH

>SGPr520_SEQID_100

PKFDPRTFHNDLALVQLWTPVSPGGSARPVCLPQEPQEPPAGTACAIAGWGALFEDGPEAEAVREARVPLLSTDTCR APPGAWPWLVRLQLGGQPLCGGVLVAASWVLTAAHCFVGCRSTRSAPNELLWTVTLAEGSRGEQAEEVPVNRILPH RALGPGLRPSTMLCAGYLAGGVDSCQGDSGGPLTCSEPGPRPREVLFGVTSWGDGCGEPGKPGVYTRVAVFKDWL QEQMSAASSSREPSCRELLAWDPPQELQADAARLCAFYARLCPGSQGACARLAHQQCLQRRRRCELRSLAHTLLGL MLLAVLLLLPLPSSWFAHGHPLYTRLPPSALQVFTLLLGAETVLGRNLDYVCEGPCGERRPSTANVTRAHGRIVGGSA

LRNAQELLGPRPGLRRLAPALALPAPALRESPLHPARELRLHSGSRAAGTRFPKRRPEPRGEANGCPGLEPLRQKLA ALQGAHAWILQVPSEHLAMNFHEVLADLGSKTLTGLFRAWVRAGLGGRHVAFSGLVGLEPATLARSLPRLLVQALQA FRVAALAEGEPEGPWMDVGQGPGLERKGHHPLNPQVPPARQP

>SGPr455_SEQID_101

PLGRTMLCAGFPDWGMDACQGDSGGPLVCRRGGGIWILAGITSWVAGCAGGSVPVRNNHVKASLGIFSKVSELMDF GPLVCRHENGPFVLYGIVSWGAGCVQPWKPGVFARVMIFLDWIQSKINGKLFSNVIKTITSFFRVGLGTVSCCSEAELE MSPDIALLYLKHKVKFGNAVQPICLPDSDDKVEPGILCLSSGWGKISKTSEYSNVLQEMELPIMDDRACNTVLKSMNLP ITQNLFTGLDRGQPLSKVGSRYITKALSSVQEVNGSQRDKIILIKFTSLDMEKQVGCDHDYVSLRSSSGVLFSKVCGKIL PSPLLAETSEAMVPFVSDTEDSGSGFELTVTAVQKSEAGSGCGSLAILVEEGTNHSAKYPDLYPSNTRCHWFICAPEK HIIKLTFEDFAVKFSPNCIYDAVVIYGDSEEKHKLAKLCGMLTITSIFSSSNMTVIYFKSDGKNRLQGFKARFTILPSESLN KFEPKLPPQNNPVSTVKAILHDVCGIPPFSPQWLSRRIAGGEEACPHCWPWQVGLRFLGDYQCGGAIINPVWILTAAH EICAVTGWGSISAELSLNVSSLDGGLASRLQQIQVHVLEREVCEHTYYSAHPGGITEKMICAGFAASGEKDFCQGDSG FMSPGPLVRVTFHALVRGAFGISYIVLKVLGPKDSKITRLSQSSNREHLVPCEDVLLTKPEGIMQIPRNSHRTTMGCQW RLVAPLNHIIQLNIINFPMKPTTFVCHGHLRVYEGFGPGKKLIGRMLMSTELSWFLSQFSTKKTTASCGETAVSMKMMY CVQLKNNPLSWTIIAGDHDRNLKESTEQVRRAKHIIVHEDFNTLSYDSDIALIQLSSPLEYNSVVRPVCLPHSAEPLFSS KPRGFFPTPRYLLDYRGRLECSWVLRVSASSMAKFTIEYLSLLGSPVCQDSVLIIYEERHSKRKTAGGLHGRRLYSMT TSIFLALQNTCYHALPHEVVLRIK

>SGPr507_2 SEQID 102

LHRGPAIPDWQRHNSHEQGRHPDLRQNLEAPVMSDRECQKTEQGKSHRNSLCVKFVKVFSRIFGEVAVATVICKDKL QTINPIQIVRYWNYSHSAPQDDLMLIKLAKPAMLNPKVQPLTLATTNVRPGTVCLLSGLDWSQENSGLWQLEPPGHLT MKYVFYLĞVLAGTFFFADSSVQKEDPAPYLVYLKSHFNPCVGVLIKPSWVLAPAHCYLPNLKVMLGNFKSRVRDGTE **QGIEVGHFMGGDVGIYTNVYKYVSWIENTAKDK**

>SGPr559 SEQID 103

KCIELIARCDGVSDCKDGEDEYRCVRVGGQNAVLQVFTAASWKTMCSDDWKGHYANVACAQLGFPSYVSSDNLRVS SLEGQFREEFVSIDHLLPDDKVTALHHSVYVREGCASGHVVTLQCTACGHRRGYSSRIVGGNMSLLSQWPWQASLQ MGENDPPAVEAPFSFRSLFGLDDLKISPVAPDADAVAAQILSLLPLKFFPIIVIGIIALILALAIGLGIHFDCSGKYRCRSSF FQGYHLCGGSVITPLWIITAAHCVYDLYLPKSWTIQVGLVSLLDNPAPSHLVEKIVYHSKYKPKRLGNDIALMKLAGPLT FNEMIQPVCLPNSEENFPDGKVCWTSGWGATEDGAGDASPVLNHAAVPLISNKICNHRDVYGGIISPSMLCAGYLTG GVDSCQGDSGGPLVCQERRLWKLVGATSFGIGCAEVNKPGVYTRVTSFLDWIHEQMERDLKT

>SGPr567 1 SEQID 104

PAQASPAQASPAQASPARASPALASLSRSSSGRSSSARSASVTTSPTRVYLVRATPVGAVPIRSSPARSAPATRATRE SPVQFWQGHTGIRYKEQRESCPKHAVRCDGVVDCKLKSDELGCVRFDWDKSLLKIYSGSSHQWLPICSSNWNDSYS **MERDSHGNASPARTPSAGASPAQASPAGTPPGRASPAQASPAQASPAGTPPGRASPAQASPAGTPPGRASPGRAS** EKTCQQLGFESAHRTTEVAHRDFANSFSILRYNSTIQESLHRSECPSQRYISLQCSHCGLRAMTGRIVGGALASDSKW PWQVSLHFGTTHICGGTLIDAQWVLTAAHCFFVTREKVLEGWKVYAGTSNLHQLPEAASIAEIIINSNYTDEEDDYDIAL MRLSKPLTLSAHIHPACLPMHGQTFSLNETCWITGFGKTRETDDKTSPFLREVQVNLIDFKKCNDYLVYDSYLTPRMM CAGDLRGGRDSCQGDSGGPLVCEQNNRWYLAGVTSWGTGCGQRNKPGVYTKVTEVLPWIYSKMESEVRFRKS

HYAGLHVCGGSILNEYWVLSAAHCFHRDKNIKIYDMYVGLVNLRVAGNHTQWYEVNRVILHPTYEMYHPIGGDVALVQ LKTRIVFSESVLPVCLATPEVNLTSANCWATGWGLVSKQGETSDELQEVQLPLILEPWCHLLYGHMSYIMPDMLCAGD >SGPr479_1_SEQID_105 MAAPASVMGPLGPSALGLLLLLLVVAPPRVAALVHRQPENQGISLTGSVACGRPSMEGKILGGVPAPERKWPWQVSV ILNAKTVCEGDSGGPLVCEFNRSWLQIGIVSWGRGCSNPLYPGVYASVSYFSKWICDNIEITPTPAQPAPALSPALGPT LSVLMAMLAGWSVL

>SGPr489 1 SEQID 106

CQGDSGGSLMCRNKKGAWDSGWSIWEAQVGGSLESRSSRPSLGNKVRLCLTNNFFKKLAGCGTWCSEQDVIVSGA EGKLHFPESLHLYYESKQRCVWTLLVPEEMHVLLSFSHLDVESCHHSYLSMYSLEDRPIGKFCGESLPSSILIGSNSLR GDCTSDYVTVHSDVERKKEIARLCGYDVPTPVLSPSSIMLISFHSDENGTCRGFQAIVSFIPKAVYPDLNISISEDESMFL MSLKMLISRNKLILLLGIVFFERGKSATLSLPKAPSCGQSLVKVQPWNYFNIFSRILGGSQVEKGSYPWQVSLKQRQKH ICGGSIVSPQWVITAAHCIANRNIVSTLNVTAGEYDLSQTDPGEQTLTIETVIIHPHFSTKKPMDYDIALLKMAGAFQFGH FVGPICLPELREQFEAGFICTTAGWGRLTEGGVLSQVLQEVNLPILTWEECVAALLTLKRPISGKTFLCTGFPDGGRDA LKFVSDATDYAAGFNLTYKALKPNYIPGCSYLTVLFEEGLIQSLNYPENYSDKANCDWIFQASKHHLIKLSFQSLEIEES

>SGPr465_1_SEQID_107

TVCGSFFQPQYPGQPSSSDYTIHEDMLCAGDLITGKAICRVNSRGPLVCPLNGTWFLMGLSSWSLDCCSPVGPRVFT HADYNELHRMGSDITLLQLHHHVEFSSHILPACLPEPTTWLAPDSSCWISGWGMVTEDVFLPEPFQLQEAEVGVMDN RWPWQASLLYLGGHICGAALIDSNWVASAAHCFQRCIFPPRAPLSTNPSDYRILLGYDQQSHPTEHSKQMTVNKIMV RLPYFTNWISQKKRESTPPDPALAPPQETPPALDSMTSQGIVHKPGLCAALLAAHMFLLLLILLGSL

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KKRKENGAVSTDKGCSQYFYAEHLSLHYPLEISAASGRLMCHFKLVAIVGYLIRLSIKSIQIEADNCVTDSLTIYDSLLPIR EYGSYNISQPCPVGSFRCSSGLCVPQAQRCDGVNDCFDESDELFCVSPQPACNTSSFRQHGPLICDGFRDCENGRD VVADVSSNNKGGLLVHFWIVFVMPRAKGHIFCEDCVAAILKDSIQTSIINRTSVGSLQGLAVDMDSVVLNGDCWSFLKK CTWKFQTSLSTLGIALKFYNYSITKKSMKGCEHGWWEINEHMYCGSYMDHQTIFRVPSPLVHIQLQCSSRLSDKPLLA VGSAYCGASVISREWLLSAAHCFHGNRLSDPTPWTAHLGMYVQGNAKFVSPVRRIVVHEYYNSQTFDYDIALLQLSIA >SGPr524_1_SEQID_108 MDKENSDVSAAPADLKISNISVQVVSAQKKLPVRRPPLPGRRLPLPGRRPPQRPIGKAKPKKQSKKKVPFWNVQNKIII EQNCTQSIPCNNRTFKCGNDICFRKQNAKCDGTVDCPDGSDEEGCTCSRSSSALHRIIGGTDTLEGGWPWQVSLHF WPETLKQLIQPICIPPTGQRVRSGEKCWVTGWGRRHEADNKGSLVLQQAEVELIDQTLCVSTYGIITSRMLCAGIMSG FTVFLFILAVIAWTLLWLYISKTESKDAFYFAGMFRITNIEFLPEYRQKESREFLSVSRTVQQVINLVYTTSAFSKFYEQS SSILYRICEPTRTLMSFVSTNNLMLVTFKSPHIRRLSGIRAYFEVIPEQKCENTVLVKDITGFEGKISSPYYPSYYPPKCK KRDACKGDSGGPLSCRRKSDGKWILTGIVSWGHGCGRPNFPGVYTRVSNFVPWIHKYVPSLL

>SGPr422 SEQID 109

DDIRQICLPEASASFQPNLTVHITGFGALYYGGESQNDLREARVKIISDDVCKQPQVYGNDIKPGMFCAGYMEGIYDAC MTLNKIKDLFAGKGQWDLAPEAEMLKPWMIAVLIVLSLTVVAVTIGLLVHFLVFDQKKEYYHGSFKILDPQINNNFGQSN TYQLKDLRETTENLVYSLKMYLSFVCHSPEEDGVKVDVIMVFQFPSTEQRAVREKKIQSILNQKIRNLRALPINASSVQV DNIHQCGATLISNTWLVTAAHCFQKYKNPHQWTVSFGTKINPPLMKRNVRRFIIHEKYRSAAREYDIAVVQVSSRVTFS NVAMVKNGNVGPGSGAGEAPGLGAGPAWSPMSSSTGELTVQASCGKRVVPLNVNRIASGVIAPKAAWPWQASLQY RGDSGGPLVTRDLKDTWYLIGIVSWGDNCGQKDKPGVYTQVTYYRNWIASKTGI

>SGPr538 SEQID 110

YDVALLRLQTALNFSDTVGAVCLPAKEQHFPKGSRCWVSGWGHTHPSHTYSSDMLQDTVVPLFSTQLCNSSCVYSG RLTHHKGVNLTDIKLNSSQEFAQLSPRLGGFLEEAWQPRNNCTSGQVVSLRCSECGARPLASRIVGGQSVAPGRWP WQASVALGFRHTCGGSVLAPRWVVTAAHCMHSFRLARLSSWRVHAGLVSHSAVRPHQGALVERIIPHPLYSAQNHD ASQPISGTLQDEEITLSCSEASAEEALLPALPKTVSFRINSEDFLLEAQVRDQPRWLLVCHEGWSPALGLQICWSLGHL ALTPRMLCAGYLDGRADACQGDSGGPLVCPDGDTWRLVGVVSWGRACAEPNHPGVYAKVAEFLDWIHDTAQDSLL MSLMLDDQPPMEAQYAEEGPGPGIFRAEPGDQQHPISQAVCWRSMRRGCAVLGALGLLAGAGVGSWLLVLYLCPA

FIG. 2R

YLGRAGASSLPQGHQVSRLVISIRLPQHLGLRPPLALLELSSRVEPSPSALPICLHPAGIPPGASCWVLGWKEPQDRVP MARHLLLPLVMLVISPIPGAFQDSALSPTQEEPEDLDCGRPEPSARIVGGSNAQPGTWPWQVSLHHGGGHICGGSLIA WPVCLPRASHRFVHGTACWATGWGDVQEADPLPLPWVLQEVELRLLGEATCQCLYSQPGPFNLTLQILPGMLCAGY QSDPQEPREENCTIALPECGKAPRPGAWPWEAQVMVPGSRPCHGALVSESWVLAPASCFLDPNSSDSPPRDLDAW PGALLEAELLGGWWCHCLYGRQGAAVPLPGDPPHALCPAYQEKEEVGSCWTHGPWISHVTRGAYLEDQLAWDWG PDGEETETQTCPPHTEHGACGLRLEAAPVGVLWPWLAEVHVAGDRVCTGILLAPGWVLAATHCVLRPGSTTVPYIEV VAAAVSILTQRICDCLYQGILPPGTLCVLYAEGQENRCEMTSAPPLLCQMTEGSWILVGMAVQGSRELFAAIGPEEAWI PEGRRDTCQGDSGGPLVCEEGGRWFQAGITSFGFGCGRRNRPGVFTAVATYEAWIREQVMGSEPGPAFPTQPQKT RVLLPSRPRAERVARLVQHENASWDNASDLALLQLRTPVNLSAASRPVCLPHPEHYFLPGSRCRLARWGRGEPALG PSWVLSAAHCFMTNGTLEPAAEWSVLLGVHSQDGPLDGAHTRAVAAIVVPANYSQVELGADLALLRLASPASLGPAV SQTVGEANFLPPSGSPHWPTGGSNLCPPELAKASGSPHAVYFLLLTLLIQS

AGTRCRVAGWGFVSDFEELPPGLMEAKVRVLDPDVCNSSWKGHLTLTMLCTRSGDSHRRGFCSADSGGPLVCRNR >SGPr542_SEQID_112 AMGLGLRGWGRPLLTVATALMLPVKPPAGSWGAQIIGGHEVTPHSRPYMASVRFGGQHHCGGFLLRARWVVSAAH CFSHRDLRTGLVVLGAHVLSTAEPTQQVFGIDALTTHPDYHPMTHANDICLLQLNGSAVLGPAVGLLRLPGRRARPPT **AHGLVSFSGLWCGDPKTPDVYTQVSAFVAWIWDVVRRSSPQPGPLPGTTRPPGEAA**

>SGPr551 SEQID 113

MLSPEVVQALLVEELLSTVNSSAAVPYRAEYEVDPEGLVILEASVKDIAALNSTLGCYRYSYVGQGQVLRLKGPDHLAS TQGQWTIQNRRLCGLRILQPYAERIPVVATAGITINFTSQISLTGPGVRVHYGLYNQSDPCPGEFLCSVNGLCVPACDG DPFVLSVQPVVFQACEVNLTLDNRLDSQGVLSTPYFPSYYSPQTHCSWHLTVPSLDYGLALWFDAYALRRQKYDLPC MPVAEAPÖVAGGÖGDGGDGEEAEPEGMFKACEDSKRKARGYLRLVPLFVLLALLVLASAGVLLWYFLGYKAEVMVS QVYSGSLRVLNRHFSQDLTRRESSAFRSETAKAQKMLKELITSTRLGTYYNSSSVYSFGEGPLTCFFWFILQIPEHRRL SCLWHLQGPKDLMLKLRLEWTLAECRDRLAMYDVAGPLEKRLITSVYGCSRQEPVVEVLASGAIMAVVWKKGLHSYY DGRPDCRDGSDEEHCDCGLQGPSSRIVGGAVSSEGEWPWQASLQVRGRHICGGALIADRWVITAAHCFQEDSMAS WITGWGALREGGPISNALQKVDVQLIPQDLCSEAYRYQVTPRMLCAGYRKGKKDACQGDSGGPLVCKALSGRWFLA IVLWTVFLGKVWQNSRWPGEVSFKVSRLLLHPYHEEDSHDYDVALLQLDHPVVRSAAVRPVCLPARSHFFEPGLHC VKDCPNGLDERNCVCRATFQCKEDSTCISLPKVCDGQPDCLNGSDEEQCQEGVPCGTFTFQCEDRSCVKKPNPQC **3LVSWGLGCGRPNYFGVYTRITGVISWIQQVVT**

>SGPr451 SEQID 114

TAPLKDVLQGSRIIGGTEAQAGAWPWVVSLQIKYGRVLVHVCGGTLVRERWVLTAAHCTKDTSDPLMWTAVIGTNNIH AEVHYISREMCNSERSYGGIIPNTSFCAGDEDGAFDTCRGDSGGPLMCYLPEYKRFFVMGITSYGHGCGRRGFPGVY DLPPSCSPASKMRLGLLSVALLFVGSSHLYSDHYSPSGRHRLGPSPEPAASSQQAEAVRKRLRRREGGAHAKDCG GRYPHTKKIKIKIKAIIIHPNFILESYVNDIALFHLKKAVRYNDYIQPICLPFDVFQILDGNTKCFISGWGRTKEEGNATNILQD IGPSFYQKWLTEHFFHASTQGILTINILRGQILIALCFVILLATT

>SGPr452_1_SEQID_115 SPPQPRTPDCRLQASLEALATLAPQPSDWLCFADLGWFEADGAAHSMGLGSSLKWAWAKPSGMPVPENDLVGIVG GHNAPPGKWPWQVSLRVYSYHWASWAHICGGSLIHPQWVLTAAHCIFWKDTDPSIYRIHAGDVYLYGGRGLLNVSRI IVHPNYVTAGLGADVALLQLPGSPLSPESLPPPYRLQQASVQVLENAVCEQPYRNASGHTGDRQLILDDMLCAGSEG RDSCYGDSGGPLVCRLRGSWRLVGVVSWGYGCTLRDFPGVYTHVQIYVLWILQQVGELP

>SGPr504 SEQID 116

IIGGHEVTPHSRPYMASVRFGGQHHCGGFLLRARWVVSAAQCFSH

>SGPr469 SEQID 117

<u> GDSGGPLVCELNGTWVQVGIVSWGIGCGRKGYPGVYTEVSFYKKWI</u>

IHVCSDTLISEEWVLTVAICFPLSPHPDFQANTSSAIAVVELPSPVSVSPVVLLICLPSSEVYLKKNTTSCWVTGWGYTGI LKSGPFRIWQGVKTKGEEGDRDTGTAGYAFTLLLLLGISGEPPEWVCGRPTVSSGIASGLGASVGQWPWQVSIRQGI >SGPr400_SEQID_118 MAGEQVTĀNVSRYPGQKTMSFPEKTFLLSYRASLLAVVTHRSNNSRGRAFESQVLPDLTAGDAADPPIPPLGPGAAL FQYIKRSYTLKELKVPLIDLQTCGDHYQNEILLHGVELIISEAMICSKLPVGQMDQCTVRIHPSGTFHRPCLPQ